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STIC-Biotech/ChemLib

110474

From: Borin, Michael  
Sent: Monday, December 15, 2003 3:56 PM  
To: STIC-Biotech/ChemLib  
Subject: Search request: 10/027038

Examiner: M. Borin  
CM1 12A01  
AU: 1631; Mailbox 12D01

Tel.: 305-4506

RE: 10/027038; peptide conjugates

Please conduct search of polypeptide SEQ ID 11 against the commercial and interference protein databases.

Thank you

Searcher: \_\_\_\_\_  
Phone: \_\_\_\_\_  
Location: \_\_\_\_\_  
Date Picked Up: 12/15/03  
Date Completed: 12/15/03  
Searcher Prep/Review: \_\_\_\_\_  
Clerical: \_\_\_\_\_  
Online time: \_\_\_\_\_

TYPE OF SEARCH:  
NA Sequences: \_\_\_\_\_  
AA Sequences: ✓  
Structures: \_\_\_\_\_  
Bibliographic: \_\_\_\_\_  
Litigation: \_\_\_\_\_  
Full text: \_\_\_\_\_  
Patent Family: \_\_\_\_\_  
Other: \_\_\_\_\_

VENDOR/COST (where applic.)  
STN: \_\_\_\_\_  
DIALOG: \_\_\_\_\_  
Questel/Orbit: \_\_\_\_\_  
DRLink: \_\_\_\_\_  
Lexis/Nexis: asp  
Sequence Sys.: \_\_\_\_\_  
WWW/Internet: \_\_\_\_\_  
Other (specify): \_\_\_\_\_



# **STIC Search Report**

## **Biotech-Chem Library**

**STIC Database Tracking Number: 110474**

**TO: Michael Borin**  
**Location: CM-1/12A01/12D01**  
**Art Unit: 1631**  
**Wednesday, December 17, 2003**

**Case Serial Number: 10/027038**

**From: Edward Hart**  
**Location: Biotech-Chem Library**  
**CM1-6B02**  
**Phone: 305-9203**

**edward.hart@uspto.gov**

### **Search Notes**

Examiner Borin,

Here are the results of the search you requested.

Please feel free to contact me if you have any questions.

Edward Hart

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 17, 2003, 16:19:10 ; Search time 22 seconds  
(without alignments)  
69.236 Million cell updates/sec

Title: US-10-027-038-11

Perfect score: 211  
Sequence: 1 MCPSPPTYPGDPGVEDLIRFYDNLQGMNCVTAAAC 36

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 328717 seqs, 42310858 residues

Total number of hits satisfying chosen parameters: 328717

Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%  
Listing first 45 summaries

Database : Issued Patente AA: \*  
1: /cgn2\_6/ptodaca/1/aa/5A COMB .pep: \*  
2: /cgn2\_6/ptodaca/1/aa/5B COMB .pep: \*  
3: /cgn2\_6/ptodaca/1/aa/6A COMB .pep: \*  
4: /cgn2\_6/ptodaca/1/aa/6B COMB .pep: \*  
5: /cgn2\_6/ptodaca/1/aa/PCTUS COMB .pep: \*  
6: /cgn2\_6/ptodaca/1/aa/Backfile1 .pep: \*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	147	69.7	36	1	US-07-776-272-18
2	86	40.8	36	1	US-07-882-923-1
3	86	40.8	36	1	US-08-264-030-1
4	86	40.8	36	1	US-08-338-395-4
5	86	40.8	36	3	US-08-307-403A-2
6	86	40.8	36	4	US-09-181-941-5
7	86	40.8	36	5	PCT-US95-14303-4
8	84	39.8	36	1	US-07-882-923-2
9	84	39.8	36	1	US-08-338-395-3
10	84	39.8	36	1	US-08-329-151-24
11	84	39.8	36	3	US-08-307-403A-1
12	84	39.8	36	5	PCT-US95-14303-3
13	84	39.8	97	3	US-09-054-393-1
14	84	39.8	97	3	US-08-594-946A-6
15	84	39.8	97	4	US-09-229-900-1
16	84	39.8	97	4	US-09-291-994-6
17	81	38.4	36	4	US-09-181-941-3
18	72	34.1	32	4	US-09-125-138-10
19	72	34.1	36	1	US-07-882-923-3
20	72	34.1	36	1	US-08-338-395-1
21	72	34.1	36	1	US-08-329-151-1
22	72	34.1	36	3	US-09-047-986B-1
23	72	34.1	36	4	US-09-181-941-4
24	72	34.1	36	5	PCT-US95-14303-1
25	72	34.1	63	4	US-09-529-727-9
26	71	33.6	36	1	US-08-329-151-9
27	68	32.2	36	1	US-08-338-395-2

28	68	32.2	36	1	US-08-329-151-2	Sequence 2, Appl1
29	68	32.2	36	3	US-09-054-393-2	Sequence 2, Appl1
30	68	32.2	36	3	US-09-047-986B-2	Sequence 2, Appl1
31	68	32.2	36	3	US-09-229-900-2	Sequence 2, Appl1
32	68	32.2	36	5	PCT-US95-14303-2	Sequence 2, Appl1
33	66	31.3	24	3	US-09-054-393-5	Sequence 5, Appl1
34	66	31.3	24	3	US-09-229-900-5	Sequence 5, Appl1
35	63	29.9	36	2	US-08-806-203-1	Sequence 1, Appl1
36	58.5	27.7	31	1	US-07-776-272-23	Sequence 23, Appl1
37	57	27.0	287	2	US-08-437-607A-2	Sequence 2, Appl1
38	54	25.6	36	4	US-09-181-941-2	Sequence 2, Appl1
39	53	25.1	694	4	US-09-433-043B-126	Sequence 126, App
40	53	25.1	752	3	US-08-975-762-61	Sequence 61, Appl
41	53	25.1	752	3	US-09-295-028-61	Sequence 61, Appl
42	53	25.1	752	4	US-09-106-583-61	Sequence 61, Appl
43	53	25.1	1404	4	US-09-345-473B-24	Sequence 24, Appl
44	52	24.6	36	4	US-09-181-941-1	Sequence 1, Appl1
45	52	24.6	228	4	US-09-336-536-11	Sequence 11, Appl1

## ALIGNMENTS

RESULT 1  
US-07-776-272-18  
Sequence 18, Application US/0776272  
Patent No. 5612454  
GENERAL INFORMATION:  
APPLICANT: Kamitama, Toshitiko  
APPLICANT: Iida, Toshii  
APPLICANT: Tajima, Masahiro  
TITLE OF INVENTION: Process for Purification of Polypeptide  
NUMBER OF SEQUENCES: 31  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Wegner, Cantor, Mueller & Player  
STREET: 1233 20th St. N.W. P.O. Box 18218  
CITY: Washington  
STATE: District of Columbia  
COUNTRY: United States of America  
ZIP: 20036-8218  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/07/776,272  
FILING DATE: 19911129  
CLASSIFICATION: 530  
ATTORNEY/AGENT INFORMATION:  
NAME: Player, William E  
REGISTRATION NUMBER: 31,409  
REFERENCE/DOCKET NUMBER: P-450-23167  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 202-887-0400  
TELEFAX: 202-887-0605  
TELEX: 440706  
INFORMATION FOR SEQ ID NO: 18:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: AMINO ACID  
TOPOLOGY: linear  
MOLECULAR TYPE: peptide  
HYPOTHETICAL: YES  
ORIGINAL SOURCE:  
ORGANISM: Avian  
US-07-776-272-18  
Query Match 69.7%, Score 147, DB 1, Length 36;  
Best Local Similarity 87.1%, Pred. No. 3.4e-12;  
Matches 27, Conservative 1, Mismatches 3, Indels 0, Gaps 0;  
QY 3 PSOPPTYPGDPGVEDLIRFYDNLQGMNCVTAAAC 33

Db 2 PSQPTPGDPAVEDLIRFYDNLQOYLNVIT 32

RESULT 2  
US-07-882-923-1

Sequence 1, Application US/07882923  
Patent No. 5328899  
GENERAL INFORMATION:  
APPLICANT: Boudlik, Jaroslav H.  
APPLICANT: Rivier, Jean R.F.  
APPLICANT: Brown, Marvin R.  
APPLICANT: Scott, Neal A.  
TITLE OF INVENTION: NPY PEPTIDE ANALOGS  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Fitch, Even, Tabin & Flannery  
STREET: 4250 Executive Square, Suite 510  
CITY: La Jolla  
STATE: CA  
COUNTRY: USA  
ZIP: 92037

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/07/882,923  
FILING DATE: 19920512  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/503,198  
FILING DATE: 30-MAR-1990  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/219,596  
FILING DATE: 15-JUL-1988  
ATTORNEY/AGENT INFORMATION:  
NAME: Schumann, James J.  
REGISTRATION NUMBER: 20,856  
REFERENCE/DOCKET NUMBER: 52864  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-1311  
TELEFAX: 619-552-0095  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: AMINO ACID  
TOPOLOGY: unknown  
MOLECULE TYPE: peptide  
US-07-882-923-1

Query Match 40.8%; Score 86; DB 1; Length 36;  
Best Local Similarity 45.2%; Pred. No. 0.00021;  
Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTPGDPAVEDLIRFYDNLQOYLNVIT 33  
Db 2 PSKPDNPGEDAPAEIDLARYYSALRYHYNLIT 32

RESULT 3  
US-08-264-030-1

Sequence 1, Application US/08264030  
Patent No. 5569742  
GENERAL INFORMATION:  
APPLICANT: KIRBY, Dean A.  
APPLICANT: RIVIER, Jean R.F.  
TITLE OF INVENTION: CENTRALLY TRUNCATED NPY CYCLIC PEPTIDE  
TITLE OF INVENTION: ANALOGS  
NUMBER OF SEQUENCES: 11  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Fitch, Even, Tabin & Flannery

STREET: 135 South La Salle Street, Suite 900  
CITY: Chicago  
STATE: IL  
COUNTRY: USA  
ZIP: 60603

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/264,030  
FILING DATE:  
CLASSIFICATION: 530  
ATTORNEY/AGENT INFORMATION:  
NAME: Schumann, James J.  
REGISTRATION NUMBER: 20,856  
REFERENCE/DOCKET NUMBER: 55649  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (619) 552-1311  
TELEFAX: (619) 552-0095  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
US-08-264-030-1

Query Match 40.8%; Score 86; DB 1; Length 36;  
Best Local Similarity 45.2%; Pred. No. 0.00021;  
Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTPGDPAVEDLIRFYDNLQOYLNVIT 33  
Db 2 PSKPDNPGEDAPAEIDLARYYSALRYHYNLIT 32

RESULT 4  
US-08-338-395-4

Sequence 4, Application US/08338395  
Patent No. 5574010  
GENERAL INFORMATION:  
APPLICANT: McFadden, David W.  
TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS WITH  
TITLE OF INVENTION: PEPTIDE YY AND ANALOGS THEREOF  
NUMBER OF SEQUENCES: 5  
CORRESPONDENCE ADDRESS:  
ADDRESSER: POMS, SMITH, LANDS & ROSE  
STREET: 2029 Century Park East 38th Floor  
CITY: Los Angeles  
STATE: CA  
COUNTRY: USA  
ZIP: 90067

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/338,395  
FILING DATE:  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: Oldenkamp, David J.  
REGISTRATION NUMBER: 29421  
REFERENCE/DOCKET NUMBER: 107012  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 310-788-5046  
TELEFAX: 310-277-1297

INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
ORIGINAL SOURCE:  
ORGANISM: PORCINE NEURAL PEPTIDE Y  
US-08-338-395-4

Query Match 40.8%; Score 86; DB 1; Length 36;  
Best Local Similarity 45.2%; Pred. No. 0.00021;  
Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTPGDPGPEVDLIRFYDNLQOMLNCVT 33  
DB 2 PSKPDNPGEDAPAEADLARYSALRHYINLIT 32

RESULT 5  
US-08-907-403A-2  
Sequence 2, Application US/08907403A  
Patent No. 6013633  
GENERAL INFORMATION:  
APPLICANT: Balaubdrumani, Ambikaipakan  
APPLICANT: Chance, William T.  
TITLE OF INVENTION: Compounds For Control  
TITLE OF INVENTION: Of Appetite, Blood Pressure, Cardiovascular  
TITLE OF INVENTION: Response, Libido, And Circadian Rhythm  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Wood, Herion & Evans, L.L.P.  
STREET: 441 Vine Street  
CITY: Cincinnati  
STATE: Ohio  
COUNTRY: USA  
ZIP: 45202-2917  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette, 3.50 inch,  
MEDIUM TYPE: 1.44 MB storage  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: Windows 95  
SOFTWARE: Microsoft Word  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/907,403A  
FILING DATE: 07-AUG-1997  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: 60/023,588  
FILING DATE: 09-AUG-1996  
ATTORNEY/AGENT INFORMATION:  
NAME: Albalany-Jenel, Stephen R.  
REGISTRATION NUMBER: 41,487  
REFERENCE/DOCKET NUMBER: UOC-113A-111  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (513) 241-2324  
TELEFAX: (513) 421-7269  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36  
TYPE: amino acid  
STRANDEDNESS:  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHEICAL: no  
ANTI-SENSE: no  
US-08-907-403A-2

Query Match 40.8%; Score 86; DB 3; Length 36;  
Best Local Similarity 45.2%; Pred. No. 0.00021;  
Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;  
QY 3 PSQPTPGDPGPEVDLIRFYDNLQOMLNCVT 33

DB 2 PSKPDNPGEDAPAEADLARYSALRHYINLIT 32

RESULT 6  
US-09-181-941-5  
Sequence 5, Application US/09181941  
Patent No. 6440690  
GENERAL INFORMATION:  
APPLICANT: Mor, Arnam  
Vouldoukis, Ioannis  
Nicolas, Pierre  
TITLE OF INVENTION: PEPTIDES FOR THE ACTIVATION  
OF THE IMMUNE SYSTEM IN HUMANS AND ANIMALS  
NUMBER OF SEQUENCES: 16  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Pennie & Edmonds LLP  
STREET: 1155 Avenue of the Americas  
CITY: New York  
STATE: NY  
COUNTRY: USA  
ZIP: 10036-2811  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: Windows  
SOFTWARE: FastSeq for Windows Version 2.0b  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/181,941  
FILING DATE: 28-Oct-1998  
CLASSIFICATION: <Unknown>  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 08/574,701  
FILING DATE: 19-DEC-1995  
APPLICATION NUMBER: FR 95 07831  
FILING DATE: 29-JUN-1995  
ATTORNEY/AGENT INFORMATION:  
NAME: Coruzzi, Laura A.  
REGISTRATION NUMBER: 30,742  
REFERENCE/DOCKET NUMBER: 3909-0021-999  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 650-493-4935  
TELEFAX: 650-493-5556  
TELEX: 66141 PENNIE  
INFORMATION FOR SEQ ID NO: 5:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: No. 6440690e  
SEQUENCE DESCRIPTION: SEQ ID NO: 5:  
US-09-181-941-5

Query Match 40.8%; Score 86; DB 4; Length 36;  
Best Local Similarity 45.2%; Pred. No. 0.00021;  
Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTPGDPGPEVDLIRFYDNLQOMLNCVT 33  
DB 2 PSKPDNPGEDAPAEADLARYSALRHYINLIT 32

RESULT 7  
PCT-US95-14303-4  
Sequence 4, Application PC/TUS9514303  
GENERAL INFORMATION:  
APPLICANT: McFadden, David W.  
TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS  
TITLE OF INVENTION: WITH PEPTIDE TY AND ANALOGS THEREOF  
NUMBER OF SEQUENCES: 5  
CORRESPONDENCE ADDRESS:  
ADDRESSER: POWS, SMITH, LANDE & ROSE

STREET: 2029 Century Park East 38th Floor  
CITY: Los Angeles  
STATE: CA  
COUNTRY: USA  
ZIP: 90067  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: PCT/US95/14303  
FILING DATE: 03 November 1995  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Oldenkamp, David J  
REGISTRATION NUMBER: 29421  
REFERENCE/DOCKET NUMBER: 107012F  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 310-788-5046  
TELEFAX: 310-277-1297  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
ORIGINAL SOURCE:  
ORGANISM: PORCINE NEURAL PEPTIDE Y  
PCT-US95-14303-4

Query Match 40.8%; Score 86; DB 5; Length 36;  
Best Local Similarity 45.2%; Pred. No. 0.00021;  
Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
DB 2 PSKPDNPGEDAPADMDARYYSALRHYNILIT 32

RESULT 8  
US-07-882-923-2  
Sequence 2, Application US/07882923  
Patent No. 5328899  
GENERAL INFORMATION:  
APPLICANT: Boudlik, Jacoblav H.  
APPLICANT: Rivier, Jean E.F.  
APPLICANT: Brown, Marvin R.  
APPLICANT: Scott, Neal A.  
TITLE OF INVENTION: NPY PEPTIDE ANALOGS  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Fitch, Even, Tabin & Flannery  
STREET: 4250 Executive Square, Suite 510  
CITY: La Jolla  
STATE: CA  
COUNTRY: USA  
ZIP: 92037  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/07/882,923  
FILING DATE: 19920512  
CLASSIFICATION: 514  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/503,198  
FILING DATE: 30-MAR-1990  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 07/219,596  
FILING DATE: 15-JUL-1988

ATTORNEY/AGENT INFORMATION:  
NAME: Schumann, James J.  
REGISTRATION NUMBER: 20,856  
REFERENCE/DOCKET NUMBER: 52864  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 619-552-1311  
TELEFAX: 619-552-0095  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: AMINO ACID  
TOPOLOGY: unknown  
MOLECULE TYPE: peptide  
US-07-882-923-2

Query Match 39.8%; Score 84; DB 1; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.00038;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
DB 2 PSKPDNPGEDAPADMDARYYSALRHYNILIT 32

RESULT 9  
US-08-338-395-3  
Sequence 3, Application US/08338395  
Patent No. 5574010  
GENERAL INFORMATION:  
APPLICANT: McFadden, David W  
TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS WITH  
NUMBER OF SEQUENCES: 5  
CORRESPONDENCE ADDRESS:  
ADDRESSER: POWS, SMITH, LANDE & ROSE  
STREET: 2029 Century Park East 38th Floor  
CITY: Los Angeles  
STATE: CA  
COUNTRY: USA  
ZIP: 90067  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/338,395  
FILING DATE:  
CLASSIFICATION: 514  
ATTORNEY/AGENT INFORMATION:  
NAME: Oldenkamp, David J  
REGISTRATION NUMBER: 29421  
REFERENCE/DOCKET NUMBER: 107012  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 310-788-5046  
TELEFAX: 310-277-1297  
INFORMATION FOR SEQ ID NO: 3:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
ORIGINAL SOURCE:  
ORGANISM: HUMAN NEUROPEPTIDE Y  
US-08-338-395-3

Query Match 39.8%; Score 84; DB 1; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.00038;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
DB 2 PSKPDNPGEDAPADMDARYYSALRHYNILIT 32

## RESULT 10

US-08-329-151-24  
Sequence 24, Application US/08329151

Patent No. 5604203

GENERAL INFORMATION:

APPLICANT: Balasubramanian, A.

TITLE OF INVENTION: ANALOGS OF PEPTIDE YY AND USES

TITLE OF INVENTION: THEREOF

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADDRESS:

ADDRESSES: Fish & Richardson

STREET: 225 Franklin Street

CITY: Boston

STATE: Massachusetts

COUNTRY: U.S.A.

ZIP: 02110-2804

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 MB

COMPUTER: IBM PS/2 Model 502 or 55SX

OPERATING SYSTEM: MS-DOS (Version 5.0)

SOFTWARE: Wordperfect (Version 5.1)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/329,151

FILING DATE:

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/038,534

FILING DATE: 3/29/93

APPLICATION NUMBER: 08/109,326

FILING DATE: 08/19/93

ATTORNEY/AGENT INFORMATION:

NAME: Paul T. Clark

REGISTRATION NUMBER: 30,162

REFERENCE/DOCKET NUMBER: 00537/105001

TELEPHONE: (617) 542-5070

TELEFAX: (617) 542-8906

TELEX: 200154

SEQUENCE CHARACTERISTICS:

SEQUENCE FOR SEQ ID NO: 24:

LENGTH: 36

TYPE: amino acid

STRANDEDNESS: N/A

TOPOLOGY: linear

FEATURE:

OTHER INFORMATION: The sequence has an amide C-terminus (i.e., CO-C

US-08-329-151-24

Query Match 39.8%; Score 84; DB 1; Length 36;

Best Local Similarity 41.9%; Pred. No. 0.00038;

Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEEDLIRFYDNLQOMLNCVT 33

Db 2 PSKPDNPGDADAPADMDARYTSALRHYINILIT 32

## RESULT 11

US-08-907-403A-1

Sequence 1, Application US/08907403A

Patent No. 6013633

GENERAL INFORMATION:

APPLICANT: Balasubramanian, Ambikaispahan

APPLICANT: Chance, William T.

TITLE OF INVENTION: Compounds For Control

TITLE OF INVENTION: Of Appetite, Blood Pressure, Cardiovascular

NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS:

ADDRESSES: Wood, Herton & Evans, L.L.P.

STREET: 441 Vine Street

CITY: Cincinnati

STATE: Ohio

COUNTRY: USA

ZIP: 45202-2917

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch,

MEDIUM TYPE: 1.44 MB storage

COMPUTER: IBM Compatible

OPERATING SYSTEM: Windows 95

SOFTWARE: Microsoft Word

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/907,403A

FILING DATE: 07-AUG-1997

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/023,588

FILING DATE: 09-AUG-1996

ATTORNEY/AGENT INFORMATION:

NAME: Albainy-Jenet, Stephen R.

REGISTRATION NUMBER: 41,487

REFERENCE/DOCKET NUMBER: UOC-113A-111

TELEPHONE: (513) 241-2324

TELEFAX: (513) 421-7269

SEQUENCE CHARACTERISTICS:

SEQUENCE FOR SEQ ID NO: 1:

LENGTH: 36

TYPE: amino acid

STRANDEDNESS:

TOPOLOGY: linear

MOLECULAR TYPE: peptide

HYDROTHERMAL: no

ANTI-SENSE: no

US-08-907-403A-1

Query Match 39.8%; Score 84; DB 3; Length 36;

Best Local Similarity 41.9%; Pred. No. 0.00038;

Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEEDLIRFYDNLQOMLNCVT 33

Db 2 PSKPDNPGDADAPADMDARYTSALRHYINILIT 32

## RESULT 12

PCT-US95-14303-3

Sequence 3, Application PC/TUS9514303

GENERAL INFORMATION:

APPLICANT: McPadden, David W

TITLE OF INVENTION: TREATMENT OF PANCREATIC TUMORS

TITLE OF INVENTION: WITH PEPTIDE YY AND ANALOGS THEREOF

NUMBER OF SEQUENCES: 5

CORRESPONDENCE ADDRESS:

ADDRESSES: POWS, SMITH, LANDR & ROSE

STREET: 2029 Century Park East 38th Floor

CITY: Los Angeles

STATE: CA

COUNTRY: USA

ZIP: 90067

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US95/14303

FILING DATE: 03 November 1995

CLASSIFICATION:

ATTORNEY/AGENT INFORMATION:

NAME: Oldenkamp, David J

REGISTRATION NUMBER: 29421

REFERENCE/DOCKET NUMBER: 107012P



TELECOMMUNICATION INFORMATION:  
TELEPHONE: 310-788-5046  
TELEFAX: 310-277-1297  
INFORMATION FOR SEQ ID NO: 3:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
ORIGINAL SOURCE:  
ORGANISM: HUMAN NEUROPEPTIDE Y  
PCT-US95-14303-3

Query Match 39.8%; Score 84; DB 5; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.00038;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQQLNCVT 33  
DB 2 PSKPDNPGEDAPADMDARYYSALRHYNILIT 32

RESULT 13  
US-09-054-393-1  
Sequence 1, Application US/09054393  
Patent No. 6017879  
GENERAL INFORMATION:  
APPLICANT: Mutter, Manfred  
APPLICANT: Lacroix, Jean S.  
APPLICANT: Grouzmann, Eric  
TITLE OF INVENTION: Template Associated NPY Y2-Receptor  
TITLE OF INVENTION: Agonists  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Vinson & Elkins LLP  
STREET: 1455 Pennsylvania Avenue, N.W.  
CITY: Washington  
STATE: D.C.  
COUNTRY: U.S.  
ZIP: 20004-1008  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/054,393  
FILING DATE:  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Sanzo, Michael A.  
REGISTRATION NUMBER: 36,912  
REFERENCE/DOCKET NUMBER: EBR350/48000  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202)639-6585  
TELEFAX: (202)639-6604  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 97 amino acids  
TYPE: amino acid  
STRANDEDNESS: not relevant  
TOPOLOGY: not relevant  
MOLECULE TYPE: peptide  
HYPOTHEICAL: NO  
ANTI-SENSE: NO  
US-09-054-393-1

Query Match 39.8%; Score 84; DB 3; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0011;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQQLNCVT 33  
DB 2 PSKPDNPGEDAPADMDARYYSALRHYNILIT 32

DB 30 PSKPDNPGEDAPADMDARYYSALRHYNILIT 60

RESULT 14  
US-08-994-946A-6  
Sequence 6, Application US/08994946A  
Patent No. 6046317  
GENERAL INFORMATION:  
APPLICANT: Koulu, Markku  
APPLICANT: Karvonen, Matti  
APPLICANT: Pesonen, Ulla-Mari  
APPLICANT: Uusitupa, Matti  
TITLE OF INVENTION: A DNA Molecule Encoding a Mutant  
TITLE OF INVENTION: Prepro-Neuropeptide Y, a Mutant Signal Peptide, and Uses  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Rothwell, Pigg, Ernst & Kurtz, P.C.  
STREET: 555 15th Street NW, Suite 701-E  
CITY: Washington  
STATE: D.C.  
COUNTRY: USA  
ZIP: 20004  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/994,946A  
FILING DATE: 19-DEC-1997  
CLASSIFICATION: 800  
ATTORNEY/AGENT INFORMATION:  
NAME: Ihnen, Jeffrey L.  
REGISTRATION NUMBER: 28,957  
REFERENCE/DOCKET NUMBER: 2328-110  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 202-783-6040  
TELEFAX: 202-783-6031  
INFORMATION FOR SEQ ID NO: 6:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 97 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-994-946A-6

Query Match 39.8%; Score 84; DB 3; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0011;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQQLNCVT 33  
DB 30 PSKPDNPGEDAPADMDARYYSALRHYNILIT 60

RESULT 15  
US-09-229-900-1  
Sequence 1, Application US/09229900  
Patent No. 6286029  
GENERAL INFORMATION:  
APPLICANT: Mutter, Manfred  
APPLICANT: Lacroix, Jean S.  
APPLICANT: Grouzmann, Eric  
TITLE OF INVENTION: Template Associated NPY Y2-Receptor  
TITLE OF INVENTION: Agonists  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Vinson & Elkins LLP  
STREET: 1455 Pennsylvania Avenue, N.W.  
CITY: Washington  
STATE: D.C.  
COUNTRY: U.S.

ZIP: 20004-1008  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/229,900  
FILING DATE:  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Sanzo, Michael A.  
REGISTRATION NUMBER: 36,912  
REFERENCE/DOCKET NUMBER: BMR50/48000  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202)639-6585  
TELEFAX: (202)639-6604  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 97 amino acids  
TYPE: amino acid  
STRANDEDNESS: not relevant  
TOPOLOGY: not relevant  
MOLECULAR TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
US-09-229-900-1

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OM protein - protein search, using sw model

Run on: December 17, 2003, 16:25:40 ; Search time 21 seconds  
(without alignments)  
92.642 Million cell updates/sec

Title: US-10-027-038-11

Perfect score: 211  
Sequence: 1 MCRSPPTGPDPEVEDLIRFDNLQGLNCVTAAAC 36

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 285895 seqs, 5404359 residues

Total number of hits satisfying chosen parameters: 285895

Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Pending Patents\_AA\_New:\*  
1: /cgn2\_6/ptodata/1/paa/PCT\_US03-18657-4  
2: /cgn2\_6/ptodata/1/paa/US06\_NEW\_COMB pep.\*  
3: /cgn2\_6/ptodata/1/paa/US07\_NEW\_COMB pep.\*  
4: /cgn2\_6/ptodata/1/paa/US08\_NEW\_COMB pep.\*  
5: /cgn2\_6/ptodata/1/paa/US09\_NEW\_COMB pep.\*  
6: /cgn2\_6/ptodata/1/paa/US10\_NEW\_COMB pep.\*  
7: /cgn2\_6/ptodata/1/paa/US60\_NEW\_COMB pep.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.	Score	Query Match	Length DB	ID	Description
1	84	39.8	36	1 PCT-US03-18657-4	Sequence 4, App11
2	84	39.8	36	5 US-09-618-361B-1	Sequence 1, App11
3	84	39.8	97	1 PCT-US03-20245-6	Sequence 6, App11
4	84	39.8	97	1 PCT-US03-20245-8	Sequence 8, App11
5	84	39.8	97	6 US-10-463-803A-1	Sequence 1, App11
6	84	39.8	97	6 US-10-686-282-7	Sequence 7, App11
7	84	39.8	97	6 US-10-686-349-7	Sequence 7, App11
8	84	39.8	36	1 PCT-US03-18657-2	Sequence 2, App11
9	84	39.8	36	6 US-10-463-803A-2	Sequence 2, App11
10	84	39.8	36	6 US-10-463-803A-19	Sequence 19, App11
11	84	39.8	97	7 US-60-490-890-1250	Sequence 1250, App11
12	84	39.8	24	6 US-10-463-803A-5	Sequence 5, App11
13	84	39.8	34	1 PCT-US03-18657-3	Sequence 5, App11
14	84	39.8	36	1 PCT-US03-18657-1	Sequence 1, App11
15	84	39.8	178	5 US-09-969-984-16	Sequence 16, App11
16	84	39.8	612	7 US-60-478-156-3222	Sequence 3222, App11
17	84	39.8	188	5 US-09-897-519A-4778	Sequence 4778, App11
18	84	39.8	537	6 US-10-322-281-322	Sequence 322, App11
19	84	39.8	1309	6 US-10-461-862-27	Sequence 27, App11
20	84	39.8	1519	6 US-10-461-862-25	Sequence 25, App11
21	84	39.8	1745	7 US-60-487-610-2383	Sequence 2383, App11
22	84	39.8	377	6 US-10-258-899A-1683	Sequence 1683, App11
23	84	39.8	406	6 US-10-258-899A-1683	Sequence 1683, App11
24	84	39.8	502	6 US-10-679-063-10187	Sequence 10187, App11
25	84	39.8	641	6 US-10-322-281-361	Sequence 361, App11
26	84	39.8	1356	7 US-60-487-610-1713	Sequence 1713, App11

27	50.5	23.9	366	6 US-10-382-000-4	Sequence 4, App11
28	50.5	23.9	542	7 US-60-485-450-1222	Sequence 1222, App11
29	50	23.7	422	6 US-10-425-114A-38524	Sequence 38524, App11
30	50	23.7	499	6 US-10-425-114A-38082	Sequence 38082, App11
31	50	23.7	560	6 US-10-425-114A-70125	Sequence 70125, App11
32	50	23.7	753	7 US-60-479-073-437	Sequence 437, App11
33	50	23.7	1266	6 US-10-367-094-85	Sequence 85, App11
34	50	23.7	1268	7 US-60-487-610-2047	Sequence 2047, App11
35	50	23.7	1281	6 US-10-367-094-87	Sequence 87, App11
36	50	23.7	1283	7 US-60-487-610-2045	Sequence 2045, App11
37	50	23.7	1679	6 US-10-367-094-89	Sequence 89, App11
38	50	23.7	1685	7 US-60-487-610-2046	Sequence 2046, App11
39	49.5	23.5	1237	1 PCT-US03-28227-4477	Sequence 4477, App11
40	49.5	23.5	1454	6 US-10-679-063-26148	Sequence 26148, App11
41	49.5	23.5	1457	6 US-10-679-063-18560	Sequence 18560, App11
42	49	23.2	20	1 PCT-US03-23875-112	Sequence 112, App11
43	49	23.2	288	5 US-09-897-516A-6710	Sequence 6710, App11
44	49	23.2	289	7 US-60-505-218-538	Sequence 538, App11
45	49	23.2	303	6 US-10-425-114A-71056	Sequence 71056, App11

## ALIGNMENTS

RESULT 1  
PCT-US03-18657-4  
Sequence 4, Application PC/TUS0318657  
GENERAL INFORMATION:  
APPLICANT: Amylin Pharmaceuticals, Inc.  
TITLE OF INVENTION: Prevention and/or Treatment of Inflammatory Bowel Disease Using  
FILE REFERENCE: 54061.8101.WO00  
CURRENT APPLICATION NUMBER: PCT/US03/18657  
CURRENT FILING DATE: 2003-06-13  
PRIOR APPLICATION NUMBER: 60/388,930  
PRIOR FILING DATE: 2002-06-14  
NUMBER OF SEQ ID NOS: 4  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 4  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: MISC FEATURE  
OTHER INFORMATION: Neuropeptide Y (NPY)  
PCT-US03-18657-4

Query Match 39.8%; Score 84; DB 1; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.0001;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;  
DB 2 PSKPDNPGEDAPEDMAYRYSALRYINILIT 32  
3 PSQPTGPDPEVEDLIRFDNLQGLNCVT 33  
US-09-618-361B-1  
Sequence 1, Application US/09618361B  
GENERAL INFORMATION:  
APPLICANT: Balaubramaniam, Ambikaipakan  
APPLICANT: Chance, William T.  
APPLICANT: University of Cincinnati  
TITLE OF INVENTION: Compounds for Control of Appetite, Blood Pressure,  
Libido and Circadian Rhythm  
FILE REFERENCE: UOC-136R  
CURRENT APPLICATION NUMBER: US/09/618,361B  
CURRENT FILING DATE: 2000-07-18  
PRIOR APPLICATION NUMBER: US/09/449,914  
PRIOR FILING DATE: 1999-12-02  
NUMBER OF SEQ ID NOS: 7  
SOFTWARE: PatentIn Version 3.1  
SEQ ID NO 1

LENGTH: 36  
TYPE: PRT  
ORGANISM: Homo sapiens  
US-09-618-361B-1

Query Match 39.8%; Score 84; DB 5; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.0001;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOQWLNCVT 33  
DB 2 PSKPDNPGEDAPADMDARYYSALRHYINLIT 32

RESULT 3  
Sequence 6, Application PC/TUS0320245  
GENERAL INFORMATION:  
APPLICANT: Qian, Su  
APPLICANT: Van der Ploeg, Leonardus, H.T.  
APPLICANT: Chen, Howard  
APPLICANT: Weingarth, Drew T.  
APPLICANT: Trumbauer, Myrna  
APPLICANT: Metzger, Joseph M.  
TITLE OF INVENTION: Agouti-related protein deficient cells,  
TITLE OF INVENTION: non-human transgenic animals and methods of selecting  
TITLE OF INVENTION: compounds which regulate energy metabolism  
FILE REFERENCE: 21033Y PCT  
CURRENT APPLICATION NUMBER: PCT/US03/20245  
CURRENT FILING DATE: 2003-06-27  
PRIOR APPLICATION NUMBER: 60/393,391  
PRIOR FILING DATE: 2002-07-03  
NUMBER OF SEQ ID NOS: 14  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 6  
LENGTH: 97  
TYPE: PRT  
ORGANISM: Mus musculus  
PCT-US03-20245-6

Query Match 39.8%; Score 84; DB 1; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0003;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOQWLNCVT 33  
DB 30 PSKPDNPGEDAPADMDARYYSALRHYINLIT 60

RESULT 4  
Sequence 8, Application PC/TUS0320245  
GENERAL INFORMATION:  
APPLICANT: Qian, Su  
APPLICANT: Van der Ploeg, Leonardus, H.T.  
APPLICANT: Chen, Howard  
APPLICANT: Weingarth, Drew T.  
APPLICANT: Trumbauer, Myrna  
APPLICANT: Metzger, Joseph M.  
TITLE OF INVENTION: Agouti-related protein deficient cells,  
TITLE OF INVENTION: non-human transgenic animals and methods of selecting  
TITLE OF INVENTION: compounds which regulate energy metabolism  
FILE REFERENCE: 21033Y PCT  
CURRENT APPLICATION NUMBER: PCT/US03/20245  
CURRENT FILING DATE: 2003-06-27  
PRIOR APPLICATION NUMBER: 60/393,391  
PRIOR FILING DATE: 2002-07-03  
NUMBER OF SEQ ID NOS: 14  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 8  
LENGTH: 97  
TYPE: PRT  
ORGANISM: Homo sapiens

PCT-US03-20245-8

Query Match 39.8%; Score 84; DB 1; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0003;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOQWLNCVT 33  
DB 30 PSKPDNPGEDAPADMDARYYSALRHYINLIT 60

RESULT 5  
US-10-463-803A-1  
Sequence 1, Application US/10463803A  
GENERAL INFORMATION:  
APPLICANT: Mutter, Manfred  
Lacroix, Jean S.  
Grouzmann, Eric  
TITLE OF INVENTION: Template Associated NPX Y2-Receptor  
Agonists  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Vinsion & Elkins LLP  
STREET: 1455 Pennsylvania Avenue, N.W.  
CITY: Washington  
STATE: D.C.  
COUNTRY: U.S.  
ZIP: 20004-1008  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/463,803A  
FILING DATE: 18-Jun-2003  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Sanzo, Michael A.  
REGISTRATION NUMBER: 36,912  
REFERENCE/DOCKET NUMBER: EMB350/48000  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202) 639-6585  
TELEFAX: (202) 639-6604  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 97 amino acids  
TYPE: amino acid  
STRANDEDNESS: not relevant  
TOPOLOGY: not relevant  
MOLECULAR TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
US-10-463-803A-1

Query Match 39.8%; Score 84; DB 6; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0003;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOQWLNCVT 33  
DB 30 PSKPDNPGEDAPADMDARYYSALRHYINLIT 60

RESULT 6  
US-10-686-282-7  
Sequence 7, Application US/10686282  
GENERAL INFORMATION:  
APPLICANT: Pfizer Inc.  
APPLICANT: Pfizer Inc.  
APPLICANT: Maw, Graham Nigel  
APPLICANT: Wayman, Christopher Peter

TITLE OF INVENTION: Compounds for the Treatment of Female Sexual Dysfunction  
FILE REFERENCE: PC10343D  
CURRENT APPLICATION NUMBER: US/10/686,282  
CURRENT FILING DATE: 2003-10-15  
PRIOR APPLICATION NUMBER: US 09/708,392  
PRIOR FILING DATE: 2000-11-08  
PRIOR APPLICATION NUMBER: US 60/175,161  
PRIOR FILING DATE: 2000-03-29  
PRIOR APPLICATION NUMBER: GB 9926437.6  
PRIOR FILING DATE: 1999-11-08  
PRIOR APPLICATION NUMBER: GB 0004021.2  
PRIOR FILING DATE: 2000-02-18  
PRIOR APPLICATION NUMBER: GB 0013001.3  
PRIOR FILING DATE: 2000-05-26  
PRIOR APPLICATION NUMBER: GB 0016563.9  
PRIOR FILING DATE: 2000-07-05  
PRIOR APPLICATION NUMBER: GB 0017141.3  
PRIOR FILING DATE: 2000-07-12  
PRIOR APPLICATION NUMBER: US 60/192,962  
PRIOR FILING DATE: 2000-03-29  
PRIOR APPLICATION NUMBER: US 60/217,479  
PRIOR FILING DATE: 2000-07-11  
PRIOR APPLICATION NUMBER: US 60/221,014  
PRIOR FILING DATE: 2000-07-27  
Remaining Prior Application data removed - See File Wrapper or PALM.  
NUMBER OF SEQ ID NOS: 20  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 7  
LENGTH: 97  
TYPE: PRT  
ORGANISM: Homo Sapiens  
US-10-686-282-7

Query Match 39.8%; Score 84; DB 6; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0003;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNIQQWLNCTV 33  
DB 30 PSKPDNFGEDAPADMDARYYSALRHYINLIT 60

RESULT 7  
US-10-686-349-7  
Sequence 7, Application US/10686349  
GENERAL INFORMATION:  
APPLICANT: Pfizer Inc.  
APPLICANT: Pfizer Limited  
APPLICANT: Maw, Graham Nigel  
APPLICANT: Wayman, Christopher Peter  
TITLE OF INVENTION: Compounds for the Treatment of Female Sexual Dysfunction  
FILE REFERENCE: PC10343C  
CURRENT APPLICATION NUMBER: US/10/686,349  
CURRENT FILING DATE: 2003-10-15  
PRIOR APPLICATION NUMBER: US 09/708,392  
PRIOR FILING DATE: 2000-11-08  
PRIOR APPLICATION NUMBER: US 60/175,161  
PRIOR FILING DATE: 2000-03-29  
PRIOR APPLICATION NUMBER: GB 9926437.6  
PRIOR FILING DATE: 1999-11-08  
PRIOR APPLICATION NUMBER: GB 0004021.2  
PRIOR FILING DATE: 2000-02-18  
PRIOR APPLICATION NUMBER: GB 0013001.3  
PRIOR FILING DATE: 2000-05-26  
PRIOR APPLICATION NUMBER: GB 0016563.9  
PRIOR FILING DATE: 2000-07-05  
PRIOR APPLICATION NUMBER: GB 0017141.3  
PRIOR FILING DATE: 2000-07-12  
PRIOR APPLICATION NUMBER: US 60/192,962  
PRIOR FILING DATE: 2000-03-29  
PRIOR APPLICATION NUMBER: US 60/217,479  
PRIOR FILING DATE: 2000-07-11  
PRIOR APPLICATION NUMBER: US 60/221,014

PRIOR FILING DATE: 2000-07-27  
Remaining Prior Application data removed - See File Wrapper or PALM.  
NUMBER OF SEQ ID NOS: 20  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 7  
LENGTH: 97  
TYPE: PRT  
ORGANISM: Homo Sapiens  
US-10-686-349-7

Query Match 39.8%; Score 84; DB 6; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.0003;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNIQQWLNCTV 33  
DB 30 PSKPDNFGEDAPADMDARYYSALRHYINLIT 60

RESULT 8  
PCT-US03-18657-2  
Sequence 2, Application PC/TUS0318657  
GENERAL INFORMATION:  
APPLICANT: Amylin Pharmaceuticals, Inc.  
TITLE OF INVENTION: Prevention and/or Treatment of Inflammatory Bowel Disease Using  
FILE REFERENCE: 54061.8101.W000  
CURRENT APPLICATION NUMBER: PCT/US03/18657  
CURRENT FILING DATE: 2003-06-13  
PRIOR APPLICATION NUMBER: 60/388,930  
PRIOR FILING DATE: 2002-06-14  
NUMBER OF SEQ ID NOS: 4  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 2  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: MISC FEATURE  
OTHER INFORMATION: Peptide YY (PY)Y  
PCT-US03-18657-2

Query Match 32.2%; Score 68; DB 1; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.015;  
Matches 13; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNIQQWLNCTV 33  
DB 2 PIKPEAFGEDASPEBLNRYYSALRHYINLIT 32

RESULT 9  
US-10-463-803A-2  
Sequence 2, Application US/10463803A  
GENERAL INFORMATION:  
APPLICANT: Mutter, Manfred  
APPLICANT: Lacroix, Jean S.  
APPLICANT: Grouzmann, Eric  
TITLE OF INVENTION: Template Associated NPV Y2-Receptor Agonists  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Vinson & Elkins LLP  
STREET: 1455 Pennsylvania Avenue, N.W.  
CITY: Washington  
STATE: D.C.  
COUNTRY: U.S.  
ZIP: 20004-1008  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/463, 803A  
FILING DATE: 18-Jun-2003  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Sanzo, Michael A.  
REGISTRATION NUMBER: 36, 912  
REFERENCE/DOCKET NUMBER: BMR350/48000  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202)639-6585  
TELEFAX: (202)639-6604  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 36 amino acids  
TYPE: amino acid  
STRANDEDNESS: not relevant  
TOPOLOGY: not relevant  
MOLECULAR TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
SEQUENCE DESCRIPTION: SEQ ID NO: 2:  
US-10-463-803A-2

Query Match 32.2% Score 68; DB 6; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.015;  
Matches 13; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
Db 2 PIKPPAPGSDASPEELNRYASLRHYLNLVT 32

RESULT 10  
US-60-500-613-19  
SEQUENCE 19, Application US/60500613  
GENERAL INFORMATION:  
APPLICANT: Immonen, Tiina  
APPLICANT: Sariola, Hannu  
APPLICANT: Saarna, Mart  
APPLICANT: Alakujala, Annina  
APPLICANT: Paeterneck, Michael  
APPLICANT: Roos, Christophe  
TITLE OF INVENTION: GDNF-RELATED NEUROPEPTIDES  
FILE REFERENCE: ICTD-0005  
CURRENT APPLICATION NUMBER: US/60/500, 613  
CURRENT FILING DATE: 2003-09-05  
NUMBER OF SEQ ID NOS: 23  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 19  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Homo sapiens  
US-60-500-613-19

Query Match 32.2% Score 68; DB 7; Length 36;  
Best Local Similarity 41.9%; Pred. No. 0.015;  
Matches 13; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
Db 2 PIKPPAPGSDASPEELNRYASLRHYLNLVT 32

RESULT 11  
US-60-490-890-1250  
SEQUENCE 1250, Application US/60490890.  
GENERAL INFORMATION:  
APPLICANT: Li, Martha  
APPLICANT: Ruppow, Brent A.  
APPLICANT: Webster, Kevin R.  
APPLICANT: Jackson, Donald  
APPLICANT: Wong, Tai W.  
TITLE OF INVENTION: BIOMARKERS OF CYCLIN-DEPENDENT KINASE MODULATION

FILE REFERENCE: D0310 PSP  
CURRENT APPLICATION NUMBER: US/60/490, 890  
CURRENT FILING DATE: 2003-07-29  
NUMBER OF SEQ ID NOS: 2779  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 1250  
LENGTH: 97  
TYPE: PRT  
ORGANISM: Homo sapiens  
US-60-490-890-1250

Query Match 32.2% Score 68; DB 7; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.044;  
Matches 13; Conservative 7; Mismatches 11; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
Db 30 PIKPPAPGSDASPEELNRYASLRHYLNLVT 60

RESULT 12  
US-10-463-803A-5  
SEQUENCE 5, Application US/10463803A  
GENERAL INFORMATION:  
APPLICANT: Mutter, Manfred  
APPLICANT: Lacroix, Jean S.  
APPLICANT: Grouzmann, Eric  
TITLE OF INVENTION: Template Associated NPY Y2-Receptor Agonists  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSER: Vinson & Elkins LLP  
STREET: 1455 Pennsylvania Avenue, N.W.  
CITY: Washington  
STATE: D.C.  
COUNTRY: U.S.  
ZIP: 20004-1008  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/10/463, 803A  
FILING DATE: 18-Jun-2003  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Sanzo, Michael A.  
REGISTRATION NUMBER: 36, 912  
REFERENCE/DOCKET NUMBER: BMR350/48000  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (202)639-6585  
TELEFAX: (202)639-6604  
INFORMATION FOR SEQ ID NO: 5:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 24 amino acids  
TYPE: amino acid  
STRANDEDNESS: not relevant  
TOPOLOGY: not relevant  
MOLECULAR TYPE: peptide  
HYPOTHETICAL: NO  
ANTI-SENSE: NO  
SEQUENCE DESCRIPTION: SEQ ID NO: 5:  
US-10-463-803A-5

Query Match 31.3% Score 66; DB 6; Length 24;  
Best Local Similarity 47.8%; Pred. No. 0.018;  
Matches 11; Conservative 4; Mismatches 8; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNL 25  
Db 2 PSKPDPGSDASPEELNRYASLRHYLNLVT 24

RESULT 13  
PCT-US03-18657-3  
Sequence 3, Application PC/TUS0318657  
GENERAL INFORMATION:  
APPLICANT: Amylin Pharmaceuticals, Inc.  
TITLE OF INVENTION: Prevention and/or Treatment of Inflammatory Bowel Disease Using  
FILE REFERENCE: 54061.8101.W000  
CURRENT APPLICATION NUMBER: PCT/US03/18657  
CURRENT FILING DATE: 2003-06-13  
PRIOR APPLICATION NUMBER: 60/388,930  
PRIOR FILING DATE: 2002-06-14  
NUMBER OF SEQ ID NOS: 4  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 3  
LENGTH: 34  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: MISC FEATURE  
OTHER INFORMATION: Peptide YY[3-36]  
PCT-US03-18657-3

Query Match 29.9%; Score 63; DB 1; Length 34;  
Best Local Similarity 41.4%; Pred. No. 0.068;  
Matches 12; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

QY 5 QPTTPGDPGPEVDLIRFYDNIQOMLNCVT 33  
DB 2 PLEPVPYGDVATPEQMAQYAADLRRYINMLT 30

RESULT 14  
PCT-US03-18657-1  
Sequence 1, Application PC/TUS0318657  
GENERAL INFORMATION:  
APPLICANT: Amylin Pharmaceuticals, Inc.  
TITLE OF INVENTION: Prevention and/or Treatment of Inflammatory Bowel Disease Using  
FILE REFERENCE: 54061.8101.W000  
CURRENT APPLICATION NUMBER: PCT/US03/18657  
CURRENT FILING DATE: 2003-06-13  
PRIOR APPLICATION NUMBER: 60/388,930  
PRIOR FILING DATE: 2002-06-14  
NUMBER OF SEQ ID NOS: 4  
SOFTWARE: PatentIn version 3.2  
SEQ ID NO 1  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: MISC FEATURE  
OTHER INFORMATION: Pancreatic polypeptide (PP)  
PCT-US03-18657-1

Query Match 29.9%; Score 63; DB 1; Length 36;  
Best Local Similarity 32.3%; Pred. No. 0.072;  
Matches 10; Conservative 10; Mismatches 11; Indels 0; Gaps 0;

QY 3 PSQPTTPGDPGPEVDLIRFYDNIQOMLNCVT 33  
DB 2 PLEPVPYGDVATPEQMAQYAADLRRYINMLT 32

RESULT 15  
US-09-969-984-16  
Sequence 16, Application US/09969984  
GENERAL INFORMATION:  
APPLICANT: INCYTE GENOMICS, INC.  
APPLICANT: TANG, Y. TOM  
APPLICANT: YUB, Henry  
APPLICANT: TAL, Preeti

APPLICANT: BURFORD, Neil  
APPLICANT: BANDMAN, Olga  
APPLICANT: BAUGHN, Mariah R.  
APPLICANT: AZIMZAI, Yaida  
APPLICANT: LU, Dzung Alma M.  
APPLICANT: PATTERSON, Chandra  
TITLE OF INVENTION: EXTRACELLULAR SIGNALING MOLECULES  
FILE REFERENCE: PE-0701-1 USA  
CURRENT APPLICATION NUMBER: US/09/969,984  
CURRENT FILING DATE: 2001-10-02  
PRIOR APPLICATION NUMBER: 60/134,949; 60/144,270; 60/146,700; 60/157,508  
PRIOR FILING DATE: 1999-05-19; 1999-07-15; 1999-07-30; 1999-10-04  
NUMBER OF SEQ ID NOS: 55  
SOFTWARE: PERL Program  
SEQ ID NO 16  
LENGTH: 178  
TYPE: PRT  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: misc feature  
OTHER INFORMATION: Incyte ID No: 5037143CD1  
US-09-969-984-16

Query Match 29.9%; Score 63; DB 5; Length 178;  
Best Local Similarity 32.3%; Pred. No. 0.41;  
Matches 10; Conservative 10; Mismatches 11; Indels 0; Gaps 0;

QY 3 PSQPTTPGDPGPEVDLIRFYDNIQOMLNCVT 33  
DB 31 PLEPVPYGDVATPEQMAQYAADLRRYINMLT 61

Search completed: December 17, 2003, 16:31:36  
Job time : 21 secs

GenCore version 5.1.6  
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OM protein - protein search, using sw model

Run on: December 17, 2003, 16:31:41 / Search time 21 Seconds  
(without alignments)

164.861 Million cell updates/sec

Title: US-10-027-038-11

Perfect score: 211  
Sequence: 1 MCPSPQTPYRGPVDELIRFYDNLQOMLNCVTAAAC 36

Scoring table: BLOSUM62  
Gapop 10.0, Gapext 0.5

Searched: 283308 seqs, 96168682 residues

Total number of hits satisfying chosen parameters: 283308

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

Database :  
1: p1r1:\*  
2: p1r2:\*  
3: p1r3:\*  
4: p1r4:\*

Pred. NO. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	143	67.8	36	2 A28578	pancreatic hormone
2	137	64.9	80	1 PCCH	pancreatic hormone
3	125	59.2	36	1 PCNO	pancreatic hormone
4	119	56.4	36	1 PCGS	pancreatic hormone
5	90	42.7	36	2 S07052	neuropeptide Y - s
6	86	40.8	36	1 NPYGY	neuropeptide Y - s
7	84	39.8	36	2 A30485	neuropeptide Y - r
8	84	39.8	36	2 B30485	neuropeptide Y - g
9	84	39.8	97	1 NRYUY	neuropeptide Y pre
10	84	39.8	97	2 A41979	neuropeptide Y pre
11	84	39.8	98	2 A25916	neuropeptide Y pre
12	82	38.9	98	2 C41979	neuropeptide Y pre
13	81	38.4	36	2 A48540	neuropeptide Y - c
14	81	38.4	36	2 A33993	neuropeptide Y - l
15	81	38.4	97	2 JCI460	neuropeptide Y pre
16	75	35.5	96	2 B41979	neuropeptide Y pre
17	72	34.1	36	1 PCXA	pancreatic peptide
18	72	34.1	36	1 PCDFY	pancreatic peptide
19	72	34.1	36	1 YYPG	peptide YY - pig
20	72	34.1	36	2 A49743	pancreatic peptide
21	72	34.1	36	2 A60416	peptide YY - dog
22	72	34.1	98	2 A29364	peptide YY precurs
23	72	34.1	104	2 I50808	pancreatic hormone
24	71	33.6	36	1 PCFG	pancreatic hormone
25	71	33.6	36	2 A26377	pancreatic peptide
26	70	33.2	36	2 A28091	pancreatic hormone
27	70	33.2	36	2 S27054	neuropeptide Y - A
28	68	32.2	36	2 A31358	peptide YY - human
29	68	32.2	90	2 S34569	peptide YY precurs

30	68	32.2	90	2 S34568	peptide YY precurs
31	68	32.2	97	2 A55914	peptide YY precurs
32	68	32.2	97	2 S33795	peptide YY (clone
33	64	30.3	36	2 J00365	pancreatic hormone
34	63	29.9	36	1 A61132	pancreatic hormone
35	63	29.9	36	1 D61132	pancreatic hormone
36	63	29.9	36	1 PCFG	pancreatic hormone
37	63	29.9	36	2 C60071	pancreatic hormone
38	63	29.9	93	1 PCDG	pancreatic hormone
39	63	29.9	95	1 PCHU	pancreatic hormone
40	62	29.4	36	1 C61132	pancreatic hormone
41	62	29.4	37	2 S26954	peptide YY-related
42	62	29.4	66	1 PCBO	pancreatic hormone
43	61	28.9	36	1 PCBO	pancreatic hormone
44	61	28.9	100	2 B28261	pancreatic hormone
45	60	28.4	36	2 B60413	pancreatic hormone

#### ALIGNMENTS

##### RESULT 1

A28578  
pancreatic hormone - ostrich  
N:Alternate names: pancreatic polypeptide  
C:Species: Struthio camelus (ostrich)  
C:Date: 19-Nov-1988 #sequence\_revision 19-Nov-1988 #text\_change 12-Apr-1995  
C:Accession: A28578  
R:Ritterhaus, D.; Oelofsen, W.  
Int. J. Pept. Protein Res. 29, 739-745, 1987  
A:Title: Purification and primary structure of ostrich pancreatic polypeptide.  
A:Reference number: A28578; MUID:87307111; PMID:3623804  
A:Accession: A28578  
A:Molecule type: protein  
A:Residues: 1-36 <lit>  
A:Note: The sequence of residues 22-23 was reported as Asn-Asp in Fig. 7 and as Asp-Asn C:Superfamily: pancreatic hormone

Query Match: 67.8%; Score 143; DB 2; Length 36;  
Best Local Similarity: 80.6%; Pred. No. 5.5e-12;  
Matches 25; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVDELIRFYDNLQOMLNCVT 33  
DB 2 PAQPTYPGDDAPVEDLIRFYDNLQOMLNCVT 32

##### RESULT 2

PCCH  
pancreatic hormone precursor - chicken  
N:Alternate names: pancreatic polypeptide precursor  
N:Contains: pancreatic hormone  
C:Species: Gallus gallus (chicken)  
C:Date: 24-Apr-1984 #sequence\_revision 03-Feb-1994 #text\_change 16-Jun-2000  
C:Accession: UN0776; A38892; A01575  
R:Nata, K.; Sugimoto, T.; Kohrt, K.; Hidaka, H.; Hattori, B.; Yamamoto, H.; Yonekura, H.  
Gene 130, 183-189, 1993  
A:Title: Structure determination and evolution of the chicken cDNA and gene encoding pre-  
A:Reference number: UN0776; MUID:93366173; PMID:8359685  
A:Accession: UN0776  
A:Molecule type: DNA  
A:Residues: 1-80 <NA>  
A:Cross-references: DDBJ:D13761; NID:G391633; PIDN:BAA02907.1; PID:G391634  
A:Accession: A38892  
A:Molecule type: mRNA  
A:Residues: 1-80 <NA>  
A:Cross-references: GB:D13760; NID:G391645; PIDN:BAA02906.1; PID:G391646  
R:Kimble, J.R.; Hayden, L.J.; Pollock, H.G.  
J. Biol. Chem. 250, 9369-9376, 1975  
A:Title: Isolation and characterization of a new pancreatic polypeptide hormone.  
A:Reference number: A01575; MUID:76069270; PMID:1194289  
A:Accession: A01575  
A:Molecule type: protein



A/Residues: 26-46, 'DN', 49-61 <KIM>  
 C/Comment: This protein acts as a regulator of pancreatic and gastrointestinal functions  
 C/Genetics:

A/Introns: 60/2

C/Superfamily: pancreatic hormone  
 C/Keywords: amidated carboxyl end; hormone; pancreas  
 F/1-25/Domain: signal sequence #status predicted <SIG>  
 F/26-61/Product: pancreatic hormone #status experimental <PCH>  
 F/61/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 64.9%; Score 137; DB 1; Length 80;  
 Best Local Similarity 80.6%; Pred. No. 8e-11;  
 Matches 25; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQOVLNCT 33  
 DB 27 PSQPTYPGDDAPVBDLIRFYDNLQOVLNVT 57

### RESULT 3

PCAO pancreatic hormone - American alligator

N/Alternate names: pancreatic polypeptide

C/Species: Alligator mississippiensis (American alligator)

C/Date: 30-Jun-1987 #sequence\_revision 30-Jun-1987 #text\_change 08-Dec-1995

C/Accession: A01577; S09341

R/Glover, I.D.; Barlow, D.J.; Plets, J.E.; Wood, S.P.; Tickle, I.J.; Blundell, T.L.; Tat

Bur, J. Biochem. 142, 379-385, 1984

A/Title: Conformational studies on the pancreatic polypeptide hormone family.

A/Reference number: A01577; PMID:84261570; PMID:6745282

A/Accession: A01577

A/Molecule type: protein

A/Residues: 1-36 <GLO>

R/ance, V.; Hamilton, J.W.; Rouse, J.B.; Kimmel, J.R.; Pollock, H.G.

Gen. Comp. Endocrinol. 55, 112-124, 1984

A/Title: Isolation and characterization of reptilian insulin, glucagon, and pancreatic p

ptide.

A/Reference number: S07210; PMID:84262419; PMID:6146554

A/Accession: S09341

A/Molecule type: protein

A/Residues: 1-21, 'N', 23-36 <LAN>

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; hormone; pancreas

F/1-36/Product: pancreatic hormone #status experimental <PCH>

F/36/Modified site: amidated carboxyl end (Phe) #status experimental

Query Match 59.2%; Score 125; DB 1; Length 36;  
 Best Local Similarity 74.2%; Pred. No. 1.2e-09;  
 Matches 23; Conservative 3; Mismatches 5; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQOVLNCT 33  
 DB 2 PLQPKYFGDAPVBDLIRFYDNLQOVLNVT 32

### RESULT 4

PCGS pancreatic hormone - goose

N/Alternate names: pancreatic polypeptide

C/Species: Anser anser (domestic goose)

C/Date: 30-Jun-1987 #sequence\_revision 30-Jun-1987 #text\_change 08-Dec-1995

C/Accession: A01576; J00006

R/Glover, I.D.; Barlow, D.J.; Plets, J.E.; Wood, S.P.; Tickle, I.J.; Blundell, T.L.; Tat

Bur, J. Biochem. 142, 379-385, 1984

A/Title: Conformational studies on the pancreatic polypeptide hormone family.

A/Reference number: A01577; PMID:84261570; PMID:6745282

A/Accession: A01576

A/Molecule type: protein

A/Residues: 1-36 <GLO>

R/Xu, Y.; Lin, N.; Zhang, Y. Sci. Sin. B Chem. Biol. Agric. Med. Earth Sci. 27, 590-592, 1984

A/Title: Isolation and sequence determination of goose pancreatic polypeptide.

A/Reference number: A94237

A/Accession: J00006

A/Molecule type: protein

A/Residues: 1-36 <XTY>

A/Note: 30-Ån was also found

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; hormone; pancreas

F/1-36/Product: pancreatic hormone #status experimental <PCH>

F/36/Modified site: amidated carboxyl end (Tyr) #status experimental

Query Match 56.4%; Score 119; DB 1; Length 36;  
 Best Local Similarity 80.8%; Pred. No. 7.3e-09;  
 Matches 21; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQOVLNCT 28  
 DB 2 PSQPTYPGDDAPVBDLIRFYDNLQOVLNCT 27

### RESULT 5

S07052 neuropeptide Y - sheep

C/Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)

C/Date: 30-Jun-1992 #sequence\_revision 30-Jun-1992 #text\_change 06-Dec-1996

C/Accession: S07052

R/Sillard, R.; Agerberth, B.; Mutt, V.; Joernvall, H.

FEBS Lett. 258, 263-265, 1989

A/Title: Sheep neuropeptide Y. A third structural type of a highly conserved peptide.

A/Reference number: S07052; PMID:90092485; PMID:2599092

A/Accession: S07052

A/Molecule type: protein

A/Residues: 1-36 <SIL>

C/Function: neuropeptide inducing a number of behavioral effects including stimulatory

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; appetite; hormone; neuropeptide

F/1-36/Product: neuropeptide Y #status experimental <MAT>

F/36/Modified site: amidated carboxyl end (Tyr) #status experimental

Query Match 42.7%; Score 90; DB 2; Length 36;  
 Best Local Similarity 48.4%; Pred. No. 4.2e-05;  
 Matches 15; Conservative 6; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQOVLNCT 33  
 DB 2 PSKPNPGDDAPVBDLIRFYDNLQOVLNCT 32

### RESULT 6

NYRGY neuropeptide Y - pig

C/Species: Sus scrofa domestica (domestic pig)

C/Date: 17-Dec-1982 #sequence\_revision 17-Dec-1982 #text\_change 06-Dec-1996

C/Accession: A01573

R/Tatemoto, K.

Proc. Natl. Acad. Sci. U.S.A. 79, 5485-5489, 1982

A/Title: Neuropeptide Y: complete amino acid sequence of the brain peptide.

A/Reference number: A01573; PMID:83039395; PMID:6957876

A/Accession: A01573

A/Molecule type: protein

A/Residues: 1-36 <YAT>

A/Note: this peptide was isolated from brain (without cerebellum and pituitary) but has

C/Function: neuropeptide inducing a number of behavioral effects including stimulatory

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; appetite; hormone; neuropeptide

F/1-36/Product: neuropeptide Y #status experimental <MAT>

F/36/Modified site: amidated carboxyl end (Tyr) #status experimental

Query Match 40.8%; Score 86; DB 1; Length 36;  
 Best Local Similarity 45.2%; Pred. No. 0.00014;  
 Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;



F/64/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 39.8%; Score 84; DB 2; Length 97;  
Best Local Similarity 41.9%; Pred. No. 0.00076;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEEDLIRFYDNLQOMLNCVT 33  
DB 30 PSKPDNPGEDAPADBDARYSALRHYNILIT 60

# RESULT 11

A25916

neuropeptide Y precursor - rat

C/Species: Rattus norvegicus (Norway rat)

C/Date: 16-Aug-1988 #sequence revision 16-Aug-1988 #text\_change 16-Jul-1999

C/Accession: A27651; A25916; A28657; A30486

R/Larhammar, D.; Ericsson, A.; Persson, H.

Proc. Natl. Acad. Sci. U.S.A. 84, 2068-2072, 1987

A/Title: Structure and expression of the rat neuropeptide Y gene.

A/Reference number: A27651, PMID:87175615, PMID:3031663

A/Accession: A27651

A/Molecule type: DNA

A/Residues: 1-98 <LAR>

A/Cross-references: GB:M15793, NID:g205759, PIDN:AAA41723.1, PID:g205761.

R.Allen, J.; Novotny, J.; Martin, J.; Heinrich, G.

Proc. Natl. Acad. Sci. U.S.A. 84, 2532-2536, 1987

A/Title: Molecular structure of mammalian neuropeptide Y: analysis by molecular cloning

A/Reference number: A25916; PMID:87175708, PMID:3031687

A/Accession: A25916

A/Molecule type: mRNA

A/Residues: 1-98 <ALL>

A/Cross-references: GB:M15880, NID:g205756; PIDN:AAA41722.1; PID:g205757

R.Higuchi, H.; Yang, H.Y.T.; Sabo, S.L.

J. Biol. Chem. 263, 6288-6295, 1988

A/Title: Rat neuropeptide Y precursor gene expression. mRNA structure, tissue distribution

A/Reference number: A28657; PMID:88198174; PMID:2834371

A/Accession: A28657

A/Molecule type: mRNA

A/Residues: 1-98 <HIG>

A/Cross-references: GB:M20373, NID:g205762; PIDN:AAA41724.1; PID:g205763

R.Cortier, R.; Galliard, R.C.; Boehlen, P.

Regul. Pept. 21, 253-261, 1988

A/Title: Isolation and sequence of rat peptide YY and neuropeptide Y.

A/Reference number: J70416; PMID:88321122; PMID:3413293

A/Accession: A30486

A/Status: preliminary

A/Molecule type: protein

A/Residues: 30-65 <COR>

A/Function: neuropeptide inducing a number of behavioral effects including stimulatory

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; appetite; hormone; neuropeptide

F/1-29/Domain: signal sequence #status predicted <SIG>

F/30-65/Product: neuropeptide Y #status experimental <MAT>

F/65-98/Domain: carboxyl-terminal propeptide #status predicted <CTP>

F/65/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 39.8%; Score 84; DB 2; Length 98;  
Best Local Similarity 41.9%; Pred. No. 0.00077;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEEDLIRFYDNLQOMLNCVT 33  
DB 31 PSKPDNPGEDAPADBDARYSALRHYNILIT 61

# RESULT 12

neuropeptide Y precursor - marbled electric ray

C/Species: Torpedo marmorata (marbled electric ray)

C/Date: 04-Mar-1993 #sequence revision 18-Nov-1994 #text\_change 16-Jul-1999

C/Accession: C41979

R.Blomqvist, A.G.; Soderberg, C.; Lundell, I.; Milner, R.J.; Larhammar, D.

Proc. Natl. Acad. Sci. U.S.A. 89, 2350-2354, 1992

A/Title: Strong evolutionary conservation of neuropeptide Y: sequences of chicken, goldf

A/Reference number: A41979; PMID:92196116, PMID:1549597

A/Accession: C41979

A/Status: preliminary; not compared with conceptual translation

A/Molecule type: nucleic acid

A/Residues: 1-98 <BLQ>

A/Cross-references: NID:M67296; NID:G213238; PIDN:AAA49281.1; PID:G213239

A/Note: sequence extracted from NCBI backbone (NCBI:88402)

A/Function: neuropeptide inducing a number of behavioral effects including stimulatory

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; appetite; hormone; neuropeptide

F/1-28/Domain: signal sequence #status predicted <SIG>

F/29-64/Product: neuropeptide Y #status predicted <MAT>

F/65-98/Domain: carboxyl-terminal propeptide #status predicted <CTP>

F/64/Modified site: amidated carboxyl end (Tyr) (amide in mature form from following gly

Query Match 38.9%; Score 82; DB 2; Length 98;  
Best Local Similarity 41.9%; Pred. No. 0.0014;  
Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEEDLIRFYDNLQOMLNCVT 33  
DB 30 PSKPDNPGEDAPADBDARYSALRHYNILIT 60

# RESULT 13

neuropeptide Y - common frog

C/Species: Rana temporaria (common frog)

C/Date: 19-Nov-1993 #sequence revision 18-Nov-1994 #text\_change 06-Dec-1996

C/Accession: A48540

R. McKay, D.M.; Shaw, C.; Halton, D.W.; Thim, L.; Buchanan, K.D.

Regul. Pept. 37, 143-153, 1992

A/Title: The primary structure and tissue distribution of an amphibian neuropeptide Y.

A/Reference number: A48540; PMID:92169199; PMID:1539111

A/Accession: A48540

A/Status: preliminary

A/Molecule type: protein

A/Residues: 1-36 <MCK>

A/Experimental source: brain

A/Note: sequence extracted from NCBI backbone (NCBI:86111)

A/Function: neuropeptide inducing a number of behavioral effects including stimulatory

C/Superfamily: pancreatic hormone

C/Keywords: amidated carboxyl end; appetite; hormone; neuropeptide

F/1-36/Product: neuropeptide Y #status experimental <MAT>

F/36/Modified site: amidated carboxyl end (Tyr) #status experimental

Query Match 38.4%; Score 81; DB 2; Length 36;  
Best Local Similarity 38.7%; Pred. No. 0.00063;  
Matches 12; Conservative 9; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEEDLIRFYDNLQOMLNCVT 33  
DB 2 PSKPDNPGEDAPADBDARYSALRHYNILIT 32

# RESULT 14

neuropeptide Y - laughing frog

C/Species: Rana ridibunda (laughing frog)

C/Date: 28-Feb-1992 #sequence revision 28-Feb-1992 #text\_change 06-Dec-1996

C/Accession: A39393

R. Chartrel, N.; Conlon, J.M.; Danger, J.M.; Fournier, A.; Tonon, M.C.; Vaudry, H.

Proc. Natl. Acad. Sci. U.S.A. 89, 3862-3866, 1992

A/Title: Characterization of melanotropin-releasing-inhibiting factor (melanostatin) from

A/Reference number: A39393; PMID:91219472; PMID:1673794

A/Accession: A39393





OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;  
 OC Gallus.  
 OC NCBI\_TaxID=9031, 9103;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC SPECIES=Chicken; TISSUE=Liver;  
 RX MEDLINE=93366173; PubMed=8359685;  
 RA Nara K., Sugimoto T., Kohri K., Hidaaka H., Hattori E., Yamamoto H.,  
 RA Yonekura H., Okamoto H.;  
 RT "Structure determination and evolution of the chicken cDNA and gene  
 RT encoding prepropancreatic polypeptide.";  
 RL Gene 130:183-189(1993).  
 RN [2]  
 RP SEQUENCE OF 26-61.  
 RC SPECIES=Chicken; PubMed=1194289;  
 RX MEDLINE=76069270; PubMed=6673760;  
 RA Kimmel J.R., Hayden L.J., Pollock H.G.;  
 RT "Isolation and characterization of a new pancreatic polypeptide  
 RT hormone.";  
 RL J. Biol. Chem. 250:9369-9376(1975).  
 RN [3]  
 RP X-RAY CRYSTALLOGRAPHY (1.4 ANGSTROMS).  
 RC SPECIES=M.galllopavo;  
 RX MEDLINE=84179397; PubMed=6673760;  
 RA Glover I., Maneef I., Pites J., Woods S., Moss D., Tickle I.,  
 RA Blundell T.L.;  
 RT "Conformational flexibility in a small globular hormone: X-ray  
 RT analysis of avian pancreatic polypeptide at 0.98-A resolution.";  
 RL Biopolymers 22:293-304(1983).  
 CC -1- FUNCTION: PANCREATIC HORMONE IS SYNTHESIZED IN PANCREATIC ISLETS  
 CC OF LANGERHANS AND ACTS AS A REGULATOR OF PANCREATIC AND  
 CC GASTROINTESTINAL FUNCTIONS.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
 CC -----  
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 CC -----  
 DR EMBL: D13761; BAA02907.1; -;  
 DR EMBL: D13760; BAA02906.1; -;  
 DR PIR: JN0776; PCCH.  
 DR PDB: 1PPT; 15-OCT-91.  
 DR InterPro: IPR001955; Pancreatic\_horm.  
 DR Pfam: PF00159; hormone3.1.  
 DR SMART: SM00309; PAH; 1.  
 DR PROSITE: PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE: PS50276; PANCREATIC\_HORMONE\_2; 1.  
 KW Hormone; Cleavage on pair of basic residues; Pancreas; Signal;  
 KW Amidation; 3D-structure.  
 FT SIGNAL 1 25 POTENTIAL.  
 FT CHAIN 26 61 PANCREATIC\_HORMONE.  
 FT PROPEP 65 80  
 FT MOD\_RES 61 61  
 FT CONFLICT 47 48 ND -> DN (IN REF. 2).  
 FT TUNN 35 36  
 FT TUNN 39 56  
 FT HELIX 39 56  
 FT TURN 57 58  
 SO SEQUENCE 80 AA; 8773 MW; 90B44E27389DB050 CRC64;

RESULT 3  
 ID PAHO\_LARAR STANDARD; PRT; 36 AA.  
 AC P41337;  
 DT 01-FEB-1995 (Rel. 31, Created)  
 DT 01-FEB-1995 (Rel. 31, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Pancreatic hormone (Pancreatic polypeptide) (Pp).  
 OS Larus argentatus (Herring gull).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Archosauria; Aves; Neognathae; Charadriiformes; Laridae; Larus.  
 OC NCBI\_TaxID=3569;  
 RN [1]  
 RP SEQUENCE.  
 RC TISSUE=Pancreas;  
 RX MEDLINE=94229519; PubMed=8174930;  
 RA Barton C.L., Shaw C., Halton D.W., Thim L.;  
 RT "Isolation and structural characterisation of herring gull (Larus  
 RT argentatus) pancreatic polypeptide.";  
 RL Gen. Comp. Endocrinol. 93:255-259(1994).  
 CC -1- FUNCTION: PANCREATIC HORMONE IS SYNTHESIZED IN PANCREATIC ISLETS  
 CC OF LANGERHANS AND ACTS AS A REGULATOR OF PANCREATIC AND  
 CC GASTROINTESTINAL FUNCTIONS.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
 DR HSSP: P01306; 1PPT.  
 DR InterPro: IPR001955; Pancreatic\_horm.  
 DR Pfam: PF00159; hormone3.1.  
 DR SMART: SM00309; PAH; 1.  
 DR PROSITE: PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE: PS50276; PANCREATIC\_HORMONE\_2; 1.  
 KW Hormone; Amidation; Pancreas.  
 FT MOD\_RES 36  
 FT MOD\_RES 36  
 SO SEQUENCE 36 AA; 4237 MW; 67831F38349C9BCE CRC64;

Query Match 61.8%; Score 130; DB 1; Length 36;  
 Best Local Similarity 74.2%; Pred. No. 4; 1e-10;  
 Matches 23; Conservative 4; Mismatches 4; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQWMLNCVT 33  
 DB 2 PVQPTYPGDPGPEVDLIRFYDNLQWMLNCVT 32

RESULT 4  
 ID PAHO\_ALIMI STANDARD; PRT; 36 AA.

AC P06305;  
 DT 01-JAN-1988 (Rel. 06, Created)  
 DT 01-JAN-1988 (Rel. 06, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Pancreatic hormone (Pancreatic polypeptide) (Pp).  
 OS Alligator mississippiensis (American alligator).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Archosauria; Crocodylia; Alligatoridae; Alligatorinae; Alligator.  
 OC NCBI\_TaxID=8496;  
 RN [1]  
 RP SEQUENCE.  
 RC MEDLINE=84261570; PubMed=6745282;  
 RA Glover I.D., Barlow D.J., Pites J.B., Wood S.P., Tickle I.J.,  
 RA Blundell T.L., Tatemoto K., Kimmel J.R., Mollmer A.,  
 RA Strassburger W., Zhang Y.S.;  
 RT "Conformational studies on the pancreatic polypeptide hormone  
 RT family.";  
 RL Eur. J. Biochem. 142:379-385(1984).  
 RN [2]  
 RP SEQUENCE.  
 RC MEDLINE=84262419; PubMed=6146554;  
 RX "Isolation and characterization of reptilian insulin, glucagon, and  
 RX pancreatic polypeptide: complete amino acid sequence of alligator  
 RX (Alligator mississippiensis) insulin and pancreatic polypeptide.";

Gen. Comp. Endocrinol. 55:112-124(1984).  
 CC -1- FUNCTION: PANCREATIC HORMONE IS SYNTHESIZED IN PANCREATIC ISLETS  
 CC OF LANGEHRMANS AND ACTS AS A REGULATOR OF PANCREATIC AND  
 CC GASTROINTESTINAL FUNCTIONS.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
 DR PIR, A01577; PCMO.  
 DR HSP, P01306; 1PPT.  
 DR InterPro, IPR001955; Pancreatic\_horm.  
 DR Pfam, PF00159; hormone3; 1.  
 DR SMART, SM00309; PAH; 1.  
 DR PROSITE, PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE, PS0276; PANCREATIC\_HORMONE\_2; 1.  
 DR Hormone; Amidation; Pancreas.  
 FT MOD RES 36  
 FT CONFLICT 22  
 FT SIGNAL 22  
 SQ SEQUENCE 36 AA; 4195 MW; F896FD2448185D9D CRC64;

Query Match 59.2%; Score 125; DB 1; Length 36;  
 Best Local Similarity 74.2%; Pred. No. 1,7e-09;  
 Matches 23; Conservative 3; Mismatches 5; Indels 0; Gaps 0;

3 PSQPTPGDGPVEDIRFYDNLQWNCCT 33  
 2 PLQPKTFGDGAPVEDLQFYDDLQOYLNVVT 32

## RESULT 5

PHO ANSAN

ID PHO ANSAN STANDARD; PRT; 36 AA.

AC P06304;

DT 01-JAN-1988 (Rel. 06, Created)

DT 01-APR-1988 (Rel. 07, Last sequence update)

DT 15-SEP-2003 (Rel. 42, Last annotation update)

DE Pancreatic hormone (pancreatic polypeptide) (pp).

DS Anser anser anser (Western graylag goose).

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

OC Archosauria; Aves; Neognathae; Anseriformes; Anatidae; Anser.

OX NCBI\_Taxid=8844;

RN [1]

RP SEQUENCE.

RX MEDLINE=85016640; PubMed=6484562;

RA Xu Y., Lin N., Zhang Y.S.;

RT "Isolation and sequence determination of goose pancreatic  
 polypeptide."

RL Sci. Sh., Ser. B, Chem. Biol. Agric. Med. Earth Sci. 27:590-592(1984).

RN [2]

RP SEQUENCE.

RX MEDLINE=84261570; PubMed=6745282;

RA Glover I.D., Bartlow D.J., Pites J.R., Wood S.P., Tickle I.J.,  
 Blundell T.L., Tatemoto K., Kimmel J.R., Mollmer A., Strassburger W.,  
 Zhang Y.S.;RT "Conformational studies on the pancreatic polypeptide hormone  
 family."

RL Eur. J. Biochem. 142:379-385(1984).

CC -1- FUNCTION: PANCREATIC HORMONE IS SYNTHESIZED IN PANCREATIC ISLETS  
 CC OF LANGEHRMANS AND ACTS AS A REGULATOR OF PANCREATIC AND  
 CC GASTROINTESTINAL FUNCTIONS.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
 DR HSP, P01306; 1PPT.  
 DR InterPro, IPR001955; Pancreatic\_horm.  
 DR Pfam, PF00159; hormone3; 1.  
 DR SMART, SM00309; PAH; 1.  
 DR PROSITE, PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE, PS0276; PANCREATIC\_HORMONE\_2; 1.  
 DR Hormone; Amidation; Pancreas.  
 FT MOD RES 36  
 FT CONFLICT 22  
 FT SIGNAL 22  
 SQ SEQUENCE 36 AA; 4391 MW; 32141F4536F427BF CRC64;

Query Match 50.2%; Score 106; DB 1; Length 36;  
 Best Local Similarity 73.1%; Pred. No. 4.1e-07;

Matches 19; Conservative 3; Mismatches 4; Indels 0; Gaps 0;

3 PSQPTPGDGPVEDIRFYDNLQWNCCT 28  
 2 PSQPTPGDGPVEDIRFYDNLQOYLNVVT 27

## RESULT 6

NEUTY SHEEP

ID NEUTY SHEEP STANDARD; PRT; 97 AA.

AC P14765; Q88PPT; Q9TS16;

DT 01-APR-1990 (Rel. 14, Created)

DT 15-SEP-2003 (Rel. 42, Last sequence update)

DT 15-SEP-2003 (Rel. 42, Last annotation update)

DE Neuropeptide Y precursor [contains: Neuropeptide Y (Neuropeptide  
 tyrosine) (NPY); C-flanking peptide of NPY (CPON)].

GN NPY.

OS Ovis aries (Sheep).

OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;

OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;

OC Bovidae; Caprinae; Ovis.

OX NCBI\_Taxid=9940;

RN [1]

RP SEQUENCE FROM N.A.

RC STRAIN=ile de France; TISSUE=Hypothalamus;

RA Pillon D., Brunet G.;

RT "Nucleotide sequence of ovine preneuropeptide Y";

RL Submitted (OCT-2001) to the EMBL/Genbank/DBD databases.

RN [2]

RP SEQUENCE OF 6-95 FROM N.A.

RA Simmons J.M., Daniel J.A., Mattern R.L., Keisler D.H.;

RL Submitted (SEP-1998) to the EMBL/Genbank/DBD databases.

RN [3]

RP SEQUENCE OF 29-64.

RX MEDLINE=90092485; PubMed=2599092;

RA Sillard R., Aguerberth B., Mutt V., Joernvall H.;

RT "Sheep neuropeptide Y. A third structural type of a highly conserved  
 peptide."

RL FEBS Lett. 258:263-265(1989).

CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN  
 CC SECRETION OF GONADOTROPIN-RELEASE HORMONE.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- TISSUE SPECIFICITY: ONE OF THE MOST ABUNDANT PEPTIDES IN THE  
 CC NERVOUS SYSTEM. ALSO FOUND IN SOME CHROMAFFIN CELLS OF THE ADRENAL  
 CC MEDULLA.  
 CC -1- SIMILARITY: Belongs to the NPY family.

CC [1]

CC [2]

CC [3]

CC [4]

CC [5]

CC [6]

CC [7]

CC [8]

CC [9]

CC [10]

CC [11]

CC [12]

CC [13]

CC [14]

CC [15]

CC [16]

CC [17]

CC [18]

CC [19]

CC [20]

CC [21]

CC [22]

CC [23]

CC [24]

CC [25]

CC [26]

CC [27]

CC [28]

CC [29]

CC [30]

CC [31]

CC [32]

CC [33]

CC [34]

CC [35]

CC [36]

CC [37]

CC [38]

CC [39]

CC [40]

CC [41]

CC [42]

CC [43]

CC [44]

CC [45]

CC [46]

CC [47]

CC [48]

EMBL, AJ41904; CAD10677.1; -  
 EMBL, AF095782; AAC6886.1; -  
 PIR, S07052; S07052.  
 DR InterPro, IPR001955; Pancreatic\_horm.  
 DR HSP, P01303; IROK.  
 DR Pfam, PF00159; hormone3; 1.  
 DR PRINTS, PR00278; PANCHORMON.  
 DR PRODOM, PD001267; Pancreatic\_horm; 1.  
 DR SMART, SM00309; PAH; 1.  
 DR PROSITE, PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE, PS0276; PANCREATIC\_HORMONE\_2; 1.  
 DR Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation.  
 FT SIGNAL 1  
 FT PEPTIDE 29  
 FT PREPIDE 68  
 FT MOD RES 64  
 FT CONFLICT 64  
 FT SIGNAL 64  
 SQ SEQUENCE 97 AA; 10750 MW; 6C2209A361CF8583 CRC64;







DR PRINTS, PRO0276; PANCROHORMONE.  
 DR PRODOM, PD001267; Pancreatic\_hormu, 1.  
 DR SMART, SM00309; PAH, 1.  
 DR PROSITB, PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITB, PS0276; PANCREATIC\_HORMONE\_2; 1.  
 DR NEUROPEPTIDE, Amidation.  
 FT MOD\_RES 36  
 SQ SEQUENCE 36 AA; 4273 MW; 0D06921202CD0DD6 CRC64;

Query Match 39.8%; Score 84; DB 1; Length 36;  
 Best Local Similarity 41.9%; Pred. No. 0.00023;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQOMLNCVT 33  
 DB 2 PSKPDNPGSDAPADMDARYSALRYHYNLT 32

RESULT 9  
 ID NEUY CHICK STANDARD; PRT; 97 AA.  
 AC P28673;  
 DT 01-DEC-1992 (Rel. 24, Created)  
 DT 01-DEC-1992 (Rel. 24, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Neuropeptide Y precursor [Contains: Neuropeptide Y (Neuropeptide tyrosine) (NPY); C-flanking peptide of NPY (CPON)].  
 GN NPY.  
 OS Gallus gallus (Chicken).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae; Gallus.  
 NCBI\_TaxID=9031;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISUB-Brain;  
 RX MEDLINE=92196116; PubMed=1549597;  
 RA Blomqvist A.G., Soederberg C., Lundell I., Milner R.J., Larhammar D.;  
 RA "Strong evolutionary conservation of neuropeptide Y: sequences of chicken, goldfish, and Torpedo marmorata DNA clones.";  
 RT Proc. Natl. Acad. Sci. U.S.A. 89:2350-2354(1992).  
 CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN SECRETION OF GONADOTROPIN-RELEASE HORMONE.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.

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CC EMBL, M87294; AAA48991.1; -  
 DR EMBL, M87295; AAA48992.1; -  
 DR EMBL, M87298; AAA48992.1; JOINED.  
 DR PIR, A1979; A1979.  
 DR HSSP, P01303; IRON.  
 DR InterPro, IPR001955; Pancreatic\_hormn.  
 DR Pfam, PF00159; hormones; 1.  
 DR PRINTS, PR00278; PANCROHORMONE.  
 DR PRODOM, PD001267; Pancreatic\_hormu, 1.  
 DR SMART, SM00309; PAH; 1.  
 DR PROSITB, PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITB, PS0276; PANCREATIC\_HORMONE\_2; 1.  
 DR Neuropeptide; Cleavage on pair of basic residues; signal; Amidation.  
 FT SIGNAL 1 28  
 FT PREPIDE 29 64  
 FT PEPTIDE 68 97  
 FT MOD\_RES 64 64  
 SQ SEQUENCE 97 AA; 11097 MW; 9C924B0829B27CB CRC64;

Query Match 39.8%; Score 84; DB 1; Length 97;  
 Best Local Similarity 41.9%; Pred. No. 0.00069;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGPEVDLIRFYDNLQOMLNCVT 33  
 DB 30 PSKPDNPGSDAPADMDARYSALRYHYNLT 60

RESULT 10  
 ID NEUY HUMAN STANDARD; PRT; 97 AA.  
 AC P01303;  
 DT 21-JUL-1986 (Rel. 01, Created)  
 DT 21-JUL-1986 (Rel. 01, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Neuropeptide Y precursor [Contains: Neuropeptide Y (Neuropeptide tyrosine) (NPY); C-flanking peptide of NPY (CPON)].  
 GN NPY.  
 OS Homo sapiens (Human).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominiidae; Homo.  
 NCBI\_TaxID=9606;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC MEDLINE=84272678; PubMed=6589611;  
 RA Minch C.D., Bloom S.R., Polak J.M., Dixon J.E.;  
 RT "Cloning, characterization, and DNA sequence of a human cDNA encoding neuropeptide tyrosine.";  
 RT Proc. Natl. Acad. Sci. U.S.A. 81:4577-4581(1984).  
 RN [2]  
 RP SEQUENCE FROM N.A.  
 RC MEDLINE=86304339; PubMed=2427515;  
 RA Minch C.D., Andrews P.C., Dixon J.E.;  
 RT "Characterization, sequence, and expression of the cloned human neuropeptide Y gene.";  
 RT J. Biol. Chem. 261:11974-11979(1986).  
 RN [3]  
 RP SEQUENCE FROM N.A.  
 RC MEDLINE=86140715; PubMed=3753985;  
 RA Takeuchi T., Gumiclo D.L., Yamada T., Weisler M.H., Minch C.D., Dixon J.E., Eddy R.E., Shows T.B.;  
 RT "Genes encoding pancreatic polypeptide and neuropeptide Y are on human chromosomes 17 and 7.";  
 RT J. Clin. Invest. 77:1038-1041(1986).  
 RN [4]  
 RP SEQUENCE FROM N.A.  
 RC Lacy M.;  
 RT Submitted (MAR-1998) to the EMBL/GenBank/DBJ databases.

-----  
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Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).

[6]

STRUCTURE BY NMR OF 29-64.

PubMed=93049324; PubMed=1425680;

RA Darbon H., Bernasau J.-M., Delenze C., Chenu J., Rousset A.,

Cambliau C.,

"solution conformation of human neuropeptide Y by 1H nuclear magnetic resonance and restrained molecular dynamics.";

RT Bur. J. Biochem. 209:765-771 (1992).

LN

[7]

STRUCTURE BY NMR OF 29-64.

PubMed=97161088; PubMed=9008359;

RA Monke S.A., Karagialis G., Howlett G.J., Norton R.S.;

RT "solution structure of human neuropeptide Y.";

J. Biomol. NMR 8:379-390 (1996).

LN

[8]

IDENTIFICATION OF CPON.

PubMed=3839058;

RA Allen J.M., Polak J.M., Bloom S.R.;

RT "Presence of the predicted C-flanking peptide of neuropeptide Y (CPON) in tissue extracts.";

LN

Neuropeptides 6:95-100 (1985).

CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN SECRETION OF GONADOTROPHIN-RELEASE HORMONE.

CC -1- SUBCELLULAR LOCATION: Secreted.

CC -1- TISSUE SPECIFICITY: ONE OF THE MOST ABUNDANT PEPTIDES IN THE NERVOUS SYSTEM. ALSO FOUND IN SOME CHROMAFFIN CELLS OF THE ADRENAL MEDULLA.

CC -1- SIMILARITY: Belongs to the NPY family.

CC

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CC

EMBL; M14298; AAA59944.1; -

DR EMBL; M14298; AAA59945.1; -

DR EMBL; M14296; AAA59945.1; JOINED.

DR EMBL; M14297; AAA59945.1; JOINED.

DR EMBL; M15789; AAA59946.1; -

DR EMBL; AC004485; -; NOT ANNOTATED CDS.

DR EMBL; BC029497; AAH29497.1; -

DR PIR; A25198; NRYU.

DR PDB; 1R0N; 17-AUG-96.

DR PDB; 1QFA; 08-APR-00.

DR Genew; HGNC:7955; NPY.

DR MIM; 162640; -

DR GO; GO:0005623; C:cell; TNS.

DR GO; GO:0005246; P:calcium channel regulator activity; TNS.

DR GO; GO:0004930; P:G-protein coupled receptor activity; TNS.

DR GO; GO:0005184; P:neuropeptide hormone activity; TNS.

DR GO; GO:0006816; P:calcium ion transport; TNS.

DR GO; GO:0006928; P:cell motility; TNS.

DR GO; GO:0008283; P:cell proliferation; TNS.

DR GO; GO:0007631; P:feeding behavior; TNS.

DR GO; GO:0007187; P:G-protein signaling, coupled to cyclic nucl. . .; TNS.

DR GO; GO:0007273; P:regulation of synapse; TNS.

DR InterPro; IPR001955; Pancreatic\_hormn.

DR Pfam; PF00159; hormone3.1.

DR PRINTS; PR00278; PANCHORMON.

DR Prodom; PD001267; Pancreatic\_hormn; 1.

DR SMART; SM00309; PAH; 1.

DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.

DR PROSITE; PS50276; PANCREATIC\_HORMONE\_2; 1.

DR Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation; 3D-structure; Polymorphism.

FT SIGNAL 1 28

FT PREPIDE 29 64 NEUROPEPTIDE Y.

FT PEPTIDE 68 97 C-FLANKING PEPTIDE OF NPY.

FT MOD\_RES 64 64 AMIDATION (G-65 PROVIDE AMIDE GROUP).

FT VARIANT 7 7 L -> P (IN dbSNP:16139).

FT VARIANT 22 22 /FTID=VAR\_014538.

FT VARIANT 53 53 L -> M (IN dbSNP:5571).

FT CONFLICT 38 39 /FTID=VAR\_014539.

FT TURN 41 44 R -> G (IN REF. 2).

FT HELIX 45 45

FT TURN 46 63

SO SEQUENCE 97 AA; 10851 MW; 832CF124321718F2 CRC64;

Query Match 39 8%; Score 84; DB 1; Length 97;

Best Local Similarity 41.9%; Pred. No. 0.00069;

Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

Db 3 PSQPTPGDGPVEDLIRFDNLCQMLNCVT 33

30 PSKEDNPGEDAPADMDARYTSALRYHYNIT 60

RESULT 11

NEUY\_MOUSE STANDARD; PRT; 97 AA.

ID PS7774; Q925V2; Q9ET27;

AC 16-OCT-2001 (Rel. 40, Created)

DT 16-OCT-2001 (Rel. 40, Last sequence update)

DT 15-SEP-2003 (Rel. 42, Last annotation update)

DE Neuropeptide Y precursor [Contains: Neuropeptide Y (Neuropeptide tyrosine) (NPY); C-flanking peptide of NPY (CPON)].

OS Mus musculus (Mouse).

OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

OX NCBI\_TaxId=10090;

LN [1]

RP SEQUENCE FROM N.A.

RC STRAIN=C57BL/6J; TISSUE=Brain;

RA MEDLINE=21085660; PubMed=11217851;

RA Kawai T., Shingawa A., Shidaka K., Yoshino M., Itoh M., Ishii Y.,

Atakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,

Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamana I.,

Salto T., Ozaki Y., Gojibori T., Bono H., Kasukawa T., Salto R.,

Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,

Pleischmann W., Gaasterland T., Giesl C., King B., Kochiwa H.,

Kuehl P., Lewis S., Matsuo Y., Nikaide I., Pesole G., Quackenbush J.,

Schirral L.M., Staudl P., Suzuki R., Tomita M., Wagner L., Washio T.,

Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,

Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,

Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,

Gastincich S., Hill D., Hofmann M., Hume D.A., Kamly M., Lee N.H.,

Lyons P., Marchionni L., Mashima J., Mazzarelli U., Mondavets P.,

Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,

Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,

Suzuki H., Toyooka K., Wang K.H., Weitz C., Whitlaker C., Wilming L.,

Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohsaki S.,

Hayashizaki Y.,

"Functional annotation of a full-length mouse cDNA collection.";

LN Nature 409:685-690 (2001).

LN [3]

SEQUENCE FROM N.A.

RP STRAIN=C57BL/6; TISSUE=Brain;

RC MEDLINE=22389257; PubMed=12477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusik K., Farmer A.A., Rubin G.M., Hong L.,

Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Schetz T.E.,

RA Brownstein M.J., Ueda T.B., Toshiyuki S., Carninci P., Prange C.,  
 RA Rana S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullaly S.J.,  
 RA Bosak S.A., McKean P.J., McKernan K.J., Malek J.A., Gamarate P.H.,  
 RA Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hallyk S.W.,  
 RA Vallatton D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,  
 RA Fahy J., Helton B., Kettman M., Madan A., Rodriguez S., Sanchez A.,  
 RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,  
 RA Blakesley R.M., Touchman J.W., Green B.D., Dickson M.C.,  
 RA Rodriguez A.C., Grimwood J., Schmitz J., Myers R.M.,  
 RA Butcherfield Y.S.N., Krzywinski M.I., Skalska U., Smalhus D.E.,  
 RA Schenck A., Schein J.E., Jones S.J.M., Marra M.A.,  
 RT "Generation and initial analysis of more than 15,000 full-length human  
 RT and mouse cDNA sequences.",  
 RL Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).  
 RN [1]  
 RP SEQUENCE OF 1-89 FROM N.A.  
 RC STRAIN=NZO, and SM/J; TISSUE=Brain;  
 RX MEDLINE=21077529; PubMed=11210195;  
 RA Taylor B.A., Wnek C., Schroeder D., Phillips S.J.;  
 RT "Multiple obesity QTLs identified in an intercross between the NZO  
 RT (New Zealand obese) and the SM (small) mouse strains.",  
 RL Mamm. Genome 12:95-103 (2001).  
 CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN  
 CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE (BY SIMILARITY).  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC TISSUE SPECIFICITY: ONE OF THE MOST ABUNDANT PEPTIDES IN THE  
 CC NERVOUS SYSTEM. ALSO FOUND IN SOME CHROMAFFIN CELLS OF THE ADRENAL  
 CC MEDULLA.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
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 CC -----  
 DR EMBL; AF273768; AAC00945.1; -  
 DR EMBL; AK029872; BAB2495.1; -  
 DR EMBL; BC043012; AAH43012.1; -  
 DR EMBL; AF286198; AAC01330.1; -  
 DR EMBL; AF286199; AAC01331.1; -  
 DR HSSP; P01303; IRON.  
 DR MGD; MG197374; NPY.  
 DR GO; GO:0001664; P:G-protein-coupled receptor binding activity; IDA.  
 DR GO; GO:0008217; P:regulation of blood pressure; IDA.  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormones; 1.  
 DR PRINTS; PR00278; PANCCHORMNE.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS00276; PANCREATIC\_HORMONE\_2; 1.  
 KW Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation.  
 FT SIGNAL; 1 28  
 FT PEPTIDE; 29 64 NEUROPEPTIDE Y.  
 FT MOD RES; 68 97 C-FLANKING PEPTIDE OF NPY.  
 FT MOD RES; 64 64 AMIDATION (G-65 PROVIDE AMIDE GROUP).  
 FT CONFLICT; 42 42 A -> R (IN REF. 4).  
 SQ SEQUENCE 97 AA; 10873 MW; 780C828FA30844B CRC64;

ID NPY\_MACMU STANDARD; PRT; 97 AA.  
 AC O9XSW6;  
 DT 15-SEP-2003 (Rel. 42, Created)  
 DT 15-SEP-2003 (Rel. 42, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Neuropeptide Y precursor [Contains: Neuropeptide Y (Neuropeptide  
 DE tyrosine) (NPY); C-flanking peptide of NPY (CPON)].  
 GN NPY.  
 OS Macaca mulatta (Rhesus macaque).  
 OC Bakayorta; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Cercopithecoidea;  
 OC Cercopithecoidea; Macaca.  
 CC NCBI\_TaxID=9544;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RA Adler L.A., Golos T.G., Terasawa E.;  
 RT "Developmental changes in NPY mRNA expression in female rhesus  
 RT monkeys.", (JUN-1999) to the EMBL/GenBank/DBJ databases.  
 CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN  
 CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE (BY SIMILARITY).  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
 CC -----  
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 CC -----  
 DR EMBL; AF162280; AAD43583.1; -  
 DR HSSP; P01303; IRON.  
 DR MGD; MG197374; NPY.  
 DR GO; GO:0001664; P:G-protein-coupled receptor binding activity; IDA.  
 DR GO; GO:0008217; P:regulation of blood pressure; IDA.  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormones; 1.  
 DR PRINTS; PR00278; PANCCHORMNE.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS00276; PANCREATIC\_HORMONE\_2; 1.  
 KW Neuropeptide; Cleavage on pair of basic residues; Signal; Amidation.  
 FT SIGNAL; 1 28  
 FT PEPTIDE; 29 64 NEUROPEPTIDE Y.  
 FT MOD RES; 68 97 C-FLANKING PEPTIDE OF NPY.  
 FT MOD RES; 64 64 AMIDATION (G-65 PROVIDE AMIDE GROUP).  
 SQ SEQUENCE 97 AA; 10840 MW; 2D2209BAC20BD55E CRC64;

Query Match 39.8%; Score 84; DB 1; Length 97;  
 Best Local Similarity 41.9%; Pred. No. 0.0069;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

Query 3 PSQPTPGDPGVEDLIRFYDNLQOMLNCVT 33  
 Db 30 PSKPDNMGEDAPADMDARYSALRHYNLIT 60

RESULT 13  
 ID NPY\_MACMU STANDARD; PRT; 98 AA.  
 AC P07808;  
 DT 01-AUG-1988 (Rel. 08, Created)  
 DT 01-AUG-1988 (Rel. 08, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Neuropeptide Y precursor [Contains: Neuropeptide Y (Neuropeptide  
 DE tyrosine) (NPY); C-flanking peptide of NPY (CPON)].  
 GN NPY.  
 OS Rattus norvegicus (Rat).  
 OC Bakayorta; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.  
 CC NCBI\_TaxID=10116;  
 RN [1]  
 RP SEQUENCE FROM N.A.

RX MEDLINE=87175708; PubMed=3031687; Heinrich G.;  
 RA Allen J., Novotny J., Martin J., Martin J.,  
 RT "Molecular structure of mammalian neuropeptide Y: analysis by  
 RT molecular cloning and computer-aided comparison with crystal  
 RT structure of avian homologue.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 84:2532-2536(1987).  
 RN [2]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE=87175615; PubMed=3031663;  
 RA Lammann D., Ericsson A., Persson H.;  
 RT "Structure and expression of the rat neuropeptide Y gene.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 84:2068-2072(1987).  
 RN [3]  
 RP SEQUENCE FROM N.A.  
 RX MEDLINE=88198174; PubMed=2834371;  
 RA Hugucl H., Yang H.-Y.T., Sabol S.L.;  
 RT "rat neuropeptide Y precursor gene expression, mRNA structure, tissue  
 RT distribution, and regulation by glucocorticoids, cyclic AMP, and  
 RT photol ester.";  
 RL J. Biol. Chem. 263:6288-6295(1988).  
 RN [4]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=AGI/seghed, BB(DR)/Mor, BN/Sanhd, DA/BKL, F344/MSHd, and  
 RA Dracheva T.V., Joe B., Hashimoto A., Dobbins D.E., Wilder R.L.,  
 RA Remmers B.F.;  
 RT "Polymorphic differences in the neuropeptide Y gene among six  
 RT autoimmune disease susceptible and resistant inbred rat strains.";  
 RL Submitted (JUN-2001) to the EMBL/Genbank/DBJ databases.  
 RN [5]  
 RP SEQUENCE OF 30-65.  
 RX MEDLINE=88321122; PubMed=3413293;  
 RA Corder R., Galliard R.C., Boehlen P.;  
 RT "Isolation and sequence of rat peptide YY and neuropeptide Y.";  
 RL Regul. Pept. 21:253-261(1988).  
 CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN  
 CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE.  
 CC -1- TISSUE SPECIFICITY: ONE OF THE MOST ABUNDANT PEPTIDES IN THE  
 CC NERVOUS SYSTEM. ALSO FOUND IN SOME CHROMAFFIN CELLS OF THE ADRENAL  
 CC MEDULLA.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
 CC -----  
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 CC -----  
 DR EMBL; M15880; AAA41722.1; -;  
 DR EMBL; M15793; AAA41723.1; JOINED.  
 DR EMBL; M15792; AAA41723.1; JOINED.  
 DR EMBL; M20373; AAA41724.1; -;  
 DR EMBL; AF392056; AAL28016.1; -;  
 DR EMBL; AF392057; AAL28017.1; -;  
 DR EMBL; AF392058; AAL28018.1; -;  
 DR EMBL; AF392059; AAL28019.1; -;  
 DR EMBL; AF392060; AAL28020.1; -;  
 DR EMBL; AF392061; AAL28021.1; -;  
 DR PIR; A27651; A25916.  
 DR HSSP; P01303; IRON.  
 DR InterPro; IPR001955; Pancreatic\_horm.  
 DR Pfam; PF00159; hormone3; 1.  
 DR PRINTS; PR00278; PANCROMONE.  
 DR PRODOM; PD001267; Pancreatic\_horm; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS0276; PANCREATIC\_HORMONE\_2; 1.  
 KW Neuropeptide; Cleavage on pair of basic residues; signal; Amidation.  
 FT SIGNAL 1 29  
 FT PEPTIDE 30 65 NEUROPEPTIDE Y.

FT PEPTIDE 69 98 C-FLANKING PEPTIDE OF NPY.  
 FT MOD RES 65 65 AMIDATION (G-66 PROVIDE AMIDE GROUP).  
 SQ SEQUENCE 98 AA; 11033 MW; E7BACER3A914B7 CRC64;  
 Query Match 39.8%; Score 84; DB 1; Length 98;  
 Best Local Similarity 41.9%; Pred. No. 0.0007;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;  
 QY 3 PSQPTYPGDPGPEVDELIRFYDNLQOMLNCVT 33  
 Db 31 PSKPDNPGEDADAEADMARYSALRYHMLIT 61  
 RESULT 14  
 ID NEWY TORMA STANDARD; PRT; 98 AA.  
 AC P28674;  
 DT 01-DEC-1992 (Rel. 24, Created)  
 DT 01-DEC-1992 (Rel. 24, Last sequence update)  
 DT 15-SEP-2003 (Rel. 42, Last annotation update)  
 DE Neuropeptide Y precursor [Contains: Neuropeptide Y (Neuropeptide  
 DE tyrosine) (NPY); C-flanking peptide of NPY (CPON)].  
 GN NPY.  
 OS Torpedo marmorata (Marbled electric ray).  
 OC Eukaryota; Metazoa; Chordata; Vertebrata; Chondrichthyes;  
 OC Elasmobranchii; Squalia; Hypnosqualia; Pristigaster; Batoidae;  
 OC Torpediniformes; Torpedinidae; Torpedo.  
 OC NCBI\_Taxid=7788;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Optic lobe;  
 RX MEDLINE=92196116; PubMed=1549597;  
 RA Blomqvist A.G., Soederberg C., Lundell I., Milner R.J.,  
 RA Lammann D.;  
 RT "Strong evolutionary conservation of neuropeptide Y: sequences of  
 RT chicken, goldfish, and Torpedo marmorata DNA clones.";  
 RL Proc. Natl. Acad. Sci. U.S.A. 89:2350-2354(1992).  
 CC -1- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN  
 CC SECRETION OF GONADOTROPHIN-RELEASE HORMONE.  
 CC -1- SUBCELLULAR LOCATION: Secreted.  
 CC -1- SIMILARITY: Belongs to the NPY family.  
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 CC -----  
 DR EMBL; M87296; AAA49281.1; -;  
 DR PIR; C41979; C41979.  
 DR HSSP; P01303; IRON.  
 DR InterPro; IPR001955; Pancreatic\_horm.  
 DR Pfam; PF00159; hormone3; 1.  
 DR PRINTS; PR00278; PANCROMONE.  
 DR PRODOM; PD001267; Pancreatic\_horm; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS0276; PANCREATIC\_HORMONE\_2; 1.  
 KW Neuropeptide; Cleavage on pair of basic residues; signal; Amidation.  
 FT SIGNAL 1 28  
 FT PEPTIDE 29 64 NEUROPEPTIDE Y.  
 FT PEPTIDE 68 98 C-FLANKING PEPTIDE OF NPY.  
 FT MOD RES 64 64 AMIDATION (G-65 PROVIDE AMIDE GROUP).  
 SQ SEQUENCE 98 AA; 11468 MW; 7959679CAD64C726 CRC64;  
 Query Match 38.9%; Score 82; DB 1; Length 98;  
 Best Local Similarity 41.9%; Pred. No. 0.0013;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;  
 QY 3 PSQPTYPGDPGPEVDELIRFYDNLQOMLNCVT 33

Db 30 PSKPDNPGEGAPADDAKYSAALHRYINLIT 60

RESULT 15

NEUZY RANRL

ID NEUZY RANRL STANDARD; PRT; 36 AA.

AC P29949;

DT 01-APR-1993 (Rel. 25, Created)

DT 01-APR-1993 (Rel. 25, Last sequence update)

DT 15-SEP-2003 (Rel. 42, Last annotation update)

DE Melanostatin (Melanotropin-release-inhibiting factor) (Neuropeptide Y)

DE (NPY).

OS Rana ridibunda (laughing frog) (Marsh frog), and

OS Rana temporaria (European common frog).

OC Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;

OC Amphibia; Batrachia; Anura; Neobatrachia; Ranoidae; Ranidae; Rana.

OX NCBI\_TaxID=8406, 8407;

RP SEQUENCE.

RC SPECIES=R. ridibunda; TISSUE=Brain;

RX MEDLINE=91219472; Pubmed=1673794;

RA Charrel N., Conlon J.M., Danger J.-M., Fournier A., Tonon M.-C.,

RA Vaudry H.;

RT "Characterization of melanotropin-release-inhibiting factor

(melanostatin) from frog brain: homology with human neuropeptide Y,"

Proc. Natl. Acad. Sci. U.S.A. 88:3862-3866(1991).

RL [2]

RN SEQUENCE.

RC SPECIES=R. temporaria; TISSUE=Brain;

RX MEDLINE=92169139; Pubmed=1539111;

RA McKay D.M., Shaw C., Halcón D.W., Thim L., Buchanan K.D.;

RT "The primary structure and tissue distribution of an amphibian

neuropeptide Y."

Regul. Pept. 37:143-153(1992).

CC -!- FUNCTION: NPY IS IMPLICATED IN THE CONTROL OF FEEDING AND IN

SECRETION OF GONADOTROPHIN-RELEASE HORMONE. NPY MAY PLAY A

PHYSIOLOGICAL ROLE IN THE REGULATION OF PITUITARY MELANOTROPHS.

CC -!- SUBCELLULAR LOCATION: Secreted.

CC -!- SIMILARITY: Belongs to the NPY family.

DR PIR; A48540; A48540.

DR HSSP; P01303; 1RCN.

DR InterPro; IPR001955; Pancreatic\_hormn.

DR Pfam; PF00159; hormone3; 1.

DR PRINTS; PR00278; PANC\_HORMONE.

DR ProDom; PD001267; Pancreatic\_hormn; 1.

DR SMART; SM00309; PAH; 1.

DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.

DR PROSITE; PS00276; PANCREATIC\_HORMONE\_2; 1.

KW Neuropeptide; Amidation.

KW MOD\_RES 36 36 AMIDATION.

FT SEQUENCE 36 AA; 4245 MW; 0D145B202CD0DD6 CRC64;

SQ

Query Match 38.4%; Score 81; DB 1; Length 36;

Best Local Similarity 38.7%; Pred. No. 0.00055;

Matches 12; Conservative 9; Mismatches 10; Indels 0; Gaps 0;

OY 3 PSOPTYFGDPGVEDLIRPDNLQWLNACT 33

DB 2 PSKPDNPGEGAPADDAKYSAALHRYINLIT 32

Search completed: December 17, 2003, 16:34:05

Job time : 32 secs

GenCore version 5.1.6  
Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - protein search, using sw model

Run on: December 17, 2003, 16:31:36 ; Search time 35 seconds  
(without alignments)  
265,426 Million cell updates/sec

Title: US-10-027-038-11

Perfect score: 211  
Sequence: 1 MCPSPPTYPGDPGPEVDLIRFYDNLQGMNCTVAAC 36

Scoring table: BLOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 830525 seqs, 258052604 residues

Total number of hits satisfying chosen parameters: 830525

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%  
Listing first 45 summaries

Database :

SPTRMBL.23.\*  
1: sp\_archaea.\*  
2: sp\_bacteria.\*  
3: sp\_fungi.\*  
4: sp\_human.\*  
5: sp\_invertebrate.\*  
6: sp\_mammal.\*  
7: sp\_mhc.\*  
8: sp\_organelle.\*  
9: sp\_phage.\*  
10: sp\_plant.\*  
11: sp\_rodent.\*  
12: sp\_virus.\*  
13: sp\_vertebrate.\*  
14: sp\_unclassified.\*  
15: sp\_virus.\*  
16: sp\_bacteriaph.\*  
17: sp\_archaeap.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	90	42.7	90	6	Q9TS16
2	90	42.7	97	6	Q8SPF7
3	86	40.8	76	6	Q9NDM5
4	84	39.8	97	6	Q9XSW6
5	82	38.9	36	13	Q9PS46
6	81	38.4	89	11	Q92SV2
7	77	36.5	99	13	Q9OWP4
8	76	36.0	96	13	Q9DGK7
9	74	35.1	36	6	Q9TR93
10	73	34.6	36	13	Q8JHE7
11	72	34.1	98	11	Q91XD0
12	67	31.8	34	6	Q9TR92
13	65	30.8	99	13	Q9OWP3
14	63	29.9	581	16	Q8DIP5
15	61.5	29.1	250	16	Q8DH44
16	60	28.4	162	16	Q9KYA3

17	59.5	28.2	436	3	Q8X0U2
18	58	27.5	81	12	Q69327
19	58	27.5	194	12	Q92584
20	58	27.5	543	12	Q40639
21	57	27.0	883	16	Q98P39
22	56.5	26.8	299	6	Q9TRT9
23	56.5	26.8	888	5	Q04135
24	56.5	26.8	888	5	Q961V7
25	56	26.5	21	13	Q9BS51
26	55.5	26.3	212	16	Q8ZQ08
27	55.5	26.3	212	16	Q8ZQ08
28	55.5	26.3	214	16	Q8ZQ08
29	55.5	26.3	365	4	Q8ZQ08
30	55	26.1	996	2	Q8ZQ08
31	55	26.1	998	2	Q8ZQ08
32	55	26.1	2911	5	Q93442
33	55	26.1	4116	4	Q8TD57
34	54.5	25.8	347	6	Q8SPS7
35	54	25.6	331	16	Q9X1N8
36	54	25.6	389	2	Q59232
37	54	25.6	799	11	Q8BNS7
38	54	25.6	1023	2	Q8KRF6
39	54	25.6	1669	11	Q9QZS0
40	54	25.6	1691	11	Q9ESQ2
41	53.5	25.4	857	15	Q8UIA1
42	53.5	25.4	857	15	Q8UIA6
43	53.5	25.4	857	15	Q8UIA0
44	53.5	25.4	857	15	Q8UIA5
45	53.5	25.4	857	15	Q8UIA4

## ALIGNMENTS

RESULT 1	ID	Q9TS16	PRELIMINARY;	PRT;	90 AA.
AC	Q9TS16;				
DT	01-MAY-2000 (TREMBLrel. 13, Created)				
DT	01-MAY-2000 (TREMBLrel. 13, Last sequence update)				
DT	01-DEC-2001 (TREMBLrel. 19, Last annotation update)				
DE	Neuropeptide Y (Fragment).				
OS	Ovis aries (Sheep).				
OC	Eukaryota; Metazoa; Chordata; Vertebrata; Euteleostomi;				
OC	Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;				
OC	Bovidae; Caprinae; Ovis.				
OX	NCBI_TaxID=9940;				
RN	[1]				
RP	SEQUENCE FROM N.A.				
RA	Simmons J.M., Daniel J.A., Mattern R.L., Keisler D.H.;				
RL	Submitted (Sep-1998) to the EMBL/Genbank/DBJ databases.				
CC	-1- SIMILARITY: BELONGS TO THE NPV / PPV / PPV / PPV FAMILY.				
DR	EMBL; AF095782; AAC6986.1; --				
DR	HSSP; P01303; IRON.				
DR	InterPro; IPR001955; Pancreatic_horm.				
DR	Pfam; PF00159; hormone3.1.				
DR	PRINTS; PR00278; PANCROMONE.				
DR	ProDom; PD001267; Pancreatic_horm.1.				
DR	SMART; SM00309; PAH; 1.				
DR	PROSITE; PS00265; PANCREATIC_HORMONE_1; 1.				
DR	PROSITE; PS00276; PANCREATIC_HORMONE_2; 1.				
KW	Amidation.				
FT	NON_TER	1			
FT	NON_TER	90			
SQ	SEQUENCE 90 AA; .9916 MW; 46FP2FB153HS5FPB CRC64;				
Query Match	42.7%; Score 90; DB 6; Length 90;				
Best Local Similarity	48.4%; Pred. No. 8.8e-05;				
Matches	15; Conservative 6; Mismatches 10; Indels 0; Gaps 0;				
QY	3 PSQPTYPGDPGPEVDLIRFYDNLQGMNCTVAAC 33				
DB	25 PSKPDNPDGADAPADLIRFYDNLQGMNCTVAAC 55				

## RESULT 2

08SPF7 PRELIMINARY; PRT; 97 AA.

AC 08SPF7, PRELIMINARY; PRT; 97 AA.  
 DT 01-JUN-2002 (TREMBLREL. 21, Created)  
 DT 01-JUN-2002 (TREMBLREL. 21, Last sequence update)  
 DT 01-MAR-2003 (TREMBLREL. 23, Last annotation update)  
 DE Neuropeptide Y precursor.

GN NPY.  
 OS Ovis aries (Sheep).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovidae;  
 OC Bovidae; Caprinae; Ovis.  
 OC NCBI\_TaxID=9940;

RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=ile de France; TISSUE=Hypothalamus;

RA Pilon D., Bruneau G.;  
 RT "Nucleotide sequence of Ovine prepro-neuropeptide Y.";  
 RL Submitted (OCT-2001) to the EMBL/Genbank/DBJ databases.

CC -1- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.  
 CC EMBL; AJ417904; CAD10677.1;  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormone3; 1.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR Amidaion; Signal.

FT SIGNAL 1 28 POTENTIAL.  
 FT CHAIN 29 64 NEUROPEPTIDE Y.  
 FT CHAIN 68 97 POTENTIAL.  
 SQ SEQUENCE 97 AA; 10750 MW; 6C2209A361CFB583 CRC64;

Query Match 42.7%; Score 90; DB 6; Length 97;  
 Best Local Similarity 48.4%; Pred. No. 9.6e-05;  
 Matches 15; Conservative 6; Mismatches 10; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 Db 30 PSKPDNPGEDAPADLARYSALRHYINLIT 60

## RESULT 3

09NOMS PRELIMINARY; PRT; 76 AA.

AC 09NOMS; PRELIMINARY; PRT; 76 AA.  
 DT 01-OCT-2000 (TREMBLREL. 15, Created)  
 DT 01-OCT-2000 (TREMBLREL. 15, Last sequence update)  
 DT 01-DEC-2001 (TREMBLREL. 19, Last annotation update)  
 DE Preneuropeptide Y (fragment).

OS Sus scrofa (Pig).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.  
 OC NCBI\_TaxID=96823;

RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Hypothalamus;

RA Matteri R.L.;  
 RT Submitted (MAY-2000) to the EMBL/Genbank/DBJ databases.

CC -1- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.  
 CC EMBL; AF264083; AAF72538.1; --

DR HSPF; P01303; IRON.  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormone3; 1.

DR PRINTS; PR00278; PANCHORMONE.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.

DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR Amidaion; Neuropeptide.

FT NON TER 1 1  
 FT CHAIN 10 >45 NEUROPEPTIDE Y.  
 FT NON TER 76 76  
 SQ SEQUENCE 76 AA; 8596 MW; 84B40EC2A4F94B2C CRC64;

Query Match 40.8%; Score 86; DB 6; Length 76;  
 Best Local Similarity 45.2%; Pred. No. 0.00026;  
 Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 Db 11 PSKPDNPGEDAPADLARYSALRHYINLIT 41

## RESULT 4

09XSM6 PRELIMINARY; PRT; 97 AA.

AC 09XSM6; PRELIMINARY; PRT; 97 AA.  
 DT 01-NOV-1999 (TREMBLREL. 12, Created)  
 DT 01-NOV-1999 (TREMBLREL. 12, Last sequence update)  
 DT 01-DEC-2001 (TREMBLREL. 19, Last annotation update)  
 DE Neuropeptide Y.

GN NPY.  
 OS Macaca mulatta (Rhesus macaque).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Mammalia; Eutheria; Primates; Catarrhini; Cercopithecoidea;  
 OC Cercopithecoidea; Macaca.  
 OC NCBI\_TaxID=9544;

RN [1]  
 RP SEQUENCE FROM N.A.  
 RA Adler L.A., Golos T.G., Terasawa E.;  
 RT "Developmental changes in NPY mRNA expression in female rhesus monkeys";  
 RL Submitted (JUN-1999) to the EMBL/Genbank/DBJ databases.

CC -1- SIMILARITY: BELONGS TO THE NPY / PPY / PYY FAMILY.  
 CC EMBL; AF162280; AAD43583.1; --

DR HSPF; P01303; IRON.  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormone3; 1.  
 DR PRINTS; PR00278; PANCHORMONE.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR Amidaion.

SQ SEQUENCE 97 AA; 10840 MW; 2D2209BAC20BD58E CRC64;

Query Match 39.8%; Score 84; DB 6; Length 97;  
 Best Local Similarity 41.9%; Pred. No. 0.00064;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

Qy 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 Db 30 PSKPDNPGEDAPADLARYSALRHYINLIT 60

## RESULT 5

09PS46 PRELIMINARY; PRT; 36 AA.

AC 09PS46; PRELIMINARY; PRT; 36 AA.  
 DT 01-MAY-2000 (TREMBLREL. 13, Created)  
 DT 01-MAY-2000 (TREMBLREL. 13, Last sequence update)  
 DT 01-OCT-2001 (TREMBLREL. 18, Last annotation update)  
 DE Neuropeptide Y, NPY=PANCREATIC polypeptide homolog.

OS Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;  
 OC Elasmobranchii; Galeomorphi; Galeidae; Carcarhiniformes;  
 OC Scyliorhinidae; Scyliorhinus.  
 OC NCBI\_TaxID=7830;

RN [1]  
 RP SEQUENCE.  
 RA MEDLINE=92396601; PubMed=1523163;  
 RX Conlon J.M., Bjerning C., Hazon N.;



RT "Structural characterization of neuropeptide Y from the brain of the  
 RT dogfish, *Scyliorhinus canicula*.";  
 RL Peptides 13:493-497(1992).  
 CC -1- SIMILARITY: BELONGS TO THE NPV / PPV / PYY FAMILY.  
 DR HSSP; P01303; IRON.  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormones; 1.  
 DR PRINTS; PR00278; PANCHEATC\_hormn; 1.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCHEATC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCHEATC\_HORMONE\_2; 1.  
 DR Amidation.  
 KW SEQUENCE. 36 AA; 4169 MW; 0D1715D0D9BD0DD6 CRC64;  
 SQ  
 Query Match 38.9%; Score 82; DB 13; Length 36;  
 Best Local Similarity 41.9%; Pred. No. 0.0004;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;  
 QY 3 PSQPTPGDPGPEVDLIRFYDNLQOMLNCVT 33  
 2 PSKPDNPGDAPADLAKYTSALRYINLIT 32  
 DB  
 RESULT 6  
 ID Q925V2 PRELIMINARY; PRT; 89 AA.  
 AC Q925V2;  
 DT 01-DEC-2001 (TREMBlrel. 19, Created)  
 DT 01-DEC-2001 (TREMBlrel. 19, Last sequence update)  
 DT 01-MAR-2003 (TREMBlrel. 23, Last annotation update)  
 DE Neuropeptide Y (Fragment).  
 OS Mus musculus (Mouse).  
 CC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 CC Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 OX NCBI\_TaxId=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=SM/J;  
 RX MEDLINE=2107529; PubMed=11210195;  
 RA Taylor B.A., Wnek C., Schroeder D., Phillips S.J.;  
 RT "Multiple obesity QTLs identified in an intercross between the NZO  
 RT (New Zealand obese) and the SM (small) mouse strains."  
 RL Mamm. Genome 12:95-103(2001).  
 CC -1- SIMILARITY: BELONGS TO THE NPV / PPV / PYY FAMILY.  
 DR EMBL; AF26198; AAG01330.1; -;  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormones; 1.  
 DR PRINTS; PR00278; PANCHEATC\_hormn; 1.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCHEATC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCHEATC\_HORMONE\_2; 1.  
 DR Amidation.  
 KW NON TER. 89  
 FT SEQUENCE 89 AA; 9943 MW; AB6052615A59D96A CRC64;  
 SQ  
 Query Match 38.4%; Score 81; DB 11; Length 89;  
 Best Local Similarity 41.9%; Pred. No. 0.0015;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;  
 QY 3 PSQPTPGDPGPEVDLIRFYDNLQOMLNCVT 33  
 30 PSKPDNPGDAPADLAKYTSALRYINLIT 60  
 DB  
 RESULT 7  
 ID Q90WF4 PRELIMINARY; PRT; 99 AA.  
 AC Q90WF4;  
 DT 01-DEC-2001 (TREMBlrel. 19, Created)  
 DT 01-DEC-2001 (TREMBlrel. 19, Last sequence update)  
 DT 01-MAR-2003 (TREMBlrel. 23, Last annotation update)

DE Neuropeptide Y.  
 GN NPV.  
 OS Parulichthys olivaceus (Plounder).  
 OC Bakaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
 OC Acanthomorpha; Acanthopterygii; Percomorpha; Pluconeciformes;  
 OC Pluconecioidei; Parulichthyidae; Parulichthys.  
 OX NCBI\_TaxId=8255;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Brain;  
 RA Kurokawa T., Suzuki T.;  
 RT "Development of neuropeptide Y related peptides in the digestive  
 RT organs during the larval stage of Japanese flounder, Parulichthys  
 RT olivaceus."  
 RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.  
 CC -1- SIMILARITY: BELONGS TO THE NPV / PPV / PYY FAMILY.  
 DR EMBL; AB055211; BAB62409.1; -;  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormones; 1.  
 DR PRINTS; PR00278; PANCHEATC\_hormn; 1.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCHEATC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCHEATC\_HORMONE\_2; 1.  
 DR Amidation.  
 KW SEQUENCE. 99 AA; 11215 MW; 6FEBD47E24CE6498 CRC64;  
 SQ  
 Query Match 36.5%; Score 77; DB 13; Length 99;  
 Best Local Similarity 38.7%; Pred. No. 0.006;  
 Matches 12; Conservative 8; Mismatches 11; Indels 0; Gaps 0;  
 QY 3 PSQPTPGDPGPEVDLIRFYDNLQOMLNCVT 33  
 30 PVKPDNPGDAPADLAKYTSALRYINLIT 60  
 DB  
 RESULT 8  
 ID Q9DCK7 PRELIMINARY; PRT; 96 AA.  
 AC Q9DCK7;  
 DT 01-MAR-2001 (TREMBlrel. 16, Created)  
 DT 01-MAR-2001 (TREMBlrel. 16, Last sequence update)  
 DT 01-DEC-2001 (TREMBlrel. 19, Last annotation update)  
 DE Neuropeptide Y. (Common carp).  
 OS Cyprinus carpio. (Common carp).  
 CC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 CC Actinopterygii; Neopterygii; Teleostei; Ostariophysi; Cypriniformes;  
 CC Cyprinidae; Cyprinus.  
 OX NCBI\_TaxId=7962;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Brain;  
 RA Yingwen L., Takeishi Y.;  
 RT "Daily rhythmic gene expression of neuropeptide Y in discrete brain of  
 RT common carp, Cyprinus carpio, under the condition of self feeding."  
 RL Submitted (JUL-2000) to the EMBL/GenBank/DBJ databases.  
 CC -1- SIMILARITY: BELONGS TO THE NPV / PPV / PYY FAMILY.  
 DR EMBL; AF287347; AAG00549.1; -;  
 DR HSSP; P01303; IRON.  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormones; 1.  
 DR PRINTS; PR00278; PANCHEATC\_hormn; 1.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCHEATC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCHEATC\_HORMONE\_2; 1.  
 DR Amidation.  
 KW SEQUENCE. 96 AA; 10987 MW; C6C5ABCD87688980 CRC64;  
 SQ  
 Query Match 36.0%; Score 76; DB 13; Length 96;  
 Best Local Similarity 35.5%; Pred. No. 0.008;  
 Matches 11; Conservative 10; Mismatches 10; Indels 0; Gaps 0;



QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 DB 30 PTKEDNPGEDAPAEELAKYTSALRYHYNLT 60

## RESULT 9

Q9TR93 PRELIMINARY, PRT, 36 AA.  
 AC Q9TR93;  
 DT 01-MAY-2000 (TREMBlrel. 13, Created)  
 DT 01-MAY-2000 (TREMBlrel. 13, Last sequence update)  
 DT 01-OCT-2001 (TREMBlrel. 18, Last annotation update)  
 DE Peptide YY, PYY(1-36).  
 OS Oryctolagus cuniculus (Rabbit).  
 OC Bkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,  
 OC Mammalia, Butheria, Lagomorpha, Leporidae, Oryctolagus.  
 OC NCBI\_TaxID=9986;  
 RN [1]  
 RP SEQUENCE.  
 RX MEDLINE=95075735; PubMed=7984499;  
 RA Grand D., Schindczek M., Struk K., Shively J., Byssesele V.E.,  
 RA Goebel H., Reeve J.R., Jr.,  
 RT "Characterization of two forms of peptide YY, PYY(1-36) and PYY(3-36),  
 in the rabbit."  
 RL Peptides 15:815-820(1994).  
 CC -1- SIMILARITY: BELONGS TO THE NPY / PYY / PYY FAMILY.  
 DR HSP, P01303; IRON  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam: PF00159; hormones; 1.  
 DR PRINTS: PR00278; PANCCHROMONE.  
 DR ProDom: PD001267; Pancreatic\_hormn; 1.  
 DR SMART: SM00309; PAH; 1.  
 DR PROSITE: PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE: PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR KAMidation.  
 KW SEQUENCE 36 AA; 4285 MW; 02D499C8086DCC64;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 DB 2 PSKPEAPGEDASPEELNRYVASLRHYLNT 32

Query Match 35.1%; Score 74; DB 6; Length 36;  
 Best Local Similarity 45.2%; Pred. No. 0.0052;  
 Matches 14; Conservative 7; Mismatches 10; Indels 0; Gaps 0;

## RESULT 10

Q8JHE7 PRELIMINARY, PRT, 36 AA.  
 AC Q8JHE7;  
 DT 01-OCT-2002 (TREMBlrel. 22, Created)  
 DT 01-OCT-2002 (TREMBlrel. 22, Last sequence update)  
 DT 01-MAR-2003 (TREMBlrel. 23, Last annotation update)  
 DE Neuropeptide Y (Fragment).  
 GN NPY.  
 OS Siniperca chuatsi (Chinese perch).  
 OC Bkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;  
 OC Actinopterygii, Neopterygii, Teleostei, Euteleostei, Neoteleostei;  
 OC Acanthomorphi, Acanthopterygii, Perciformes, Percoidae;  
 OC Siniperacidae, Siniperca.  
 OC NCBI\_TaxID=119488;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RA Liang X., Bai J., Lao H.,  
 RA "Mandarin fish (Siniperca chuatsi) NPY mature peptide."  
 RL Submitted (MAY-2002) to the EMBL/GenBank/DBJ databases.  
 DR EMBL: AF514858; AAM51821.1;  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam: PF00159; hormones; 1.  
 DR PRINTS: PR00278; PANCCHROMONE.  
 DR ProDom: PD001267; Pancreatic\_hormn; 1.  
 DR SMART: SM00309; PAH; 1.

DR PROSITE: PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE: PS50276; PANCREATIC\_HORMONE\_2; 1.  
 FT NON TER 1 1  
 FT NON TER 36 36  
 SQ SEQUENCE 36 AA; 4267 MW; 17A32293A8667CC6 CRC64;

Query Match 34.6%; Score 73; DB 13; Length 36;  
 Best Local Similarity 35.5%; Pred. No. 0.0071;  
 Matches 11; Conservative 9; Mismatches 11; Indels 0; Gaps 0;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 DB 2 PVKPEAPGEDASPEELNRYVASLRHYLNT 32

## RESULT 11

Q91XD0 PRELIMINARY, PRT, 98 AA.  
 ID Q91XD0;  
 AC Q91XD0;  
 DT 01-DEC-2001 (TREMBlrel. 19, Created)  
 DT 01-DEC-2001 (TREMBlrel. 19, Last sequence update)  
 DT 01-MAR-2003 (TREMBlrel. 23, Last annotation update)  
 DE Unknown (Protein for MGC19143).  
 OS Mus musculus (Mouse).  
 OC Bkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;  
 OC Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.  
 OC NCBI\_TaxID=10090;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Colon;  
 RA Strauberg R.,  
 RL Submitted (JUL-2001) to the EMBL/GenBank/DBJ databases.  
 CC EMBL: BC010821; AAH10821.1;  
 DR EMBL: BC010821; AAH10821.1;  
 DR InterPro: IPR001955; Pancreatic\_hormn.  
 DR Pfam: PF00159; hormones; 1.  
 DR PRINTS: PR00278; PANCCHROMONE.  
 DR ProDom: PD001267; Pancreatic\_hormn; 1.  
 DR SMART: SM00309; PAH; 1.  
 DR PROSITE: PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE: PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR KAMidation.  
 KW SEQUENCE 98 AA; 11064 MW; 7AF165A1052C3249 CRC64;

QY 3 PSQPTYPGDPGVEDLIRFYDNLQOMLNCVT 33  
 DB 30 PAKPEAPGEDASPEELNRYVASLRHYLNT 60

Query Match 34.1%; Score 72; DB 11; Length 98;  
 Best Local Similarity 41.9%; Pred. No. 0.029;  
 Matches 13; Conservative 8; Mismatches 10; Indels 0; Gaps 0;

## RESULT 12

Q9TR92 PRELIMINARY, PRT, 34 AA.  
 ID Q9TR92;  
 AC Q9TR92;  
 DT 01-MAY-2000 (TREMBlrel. 13, Created)  
 DT 01-MAY-2000 (TREMBlrel. 13, Last sequence update)  
 DT 01-OCT-2001 (TREMBlrel. 18, Last annotation update)  
 DE Peptide YY, PYY(3-36).  
 OS Oryctolagus cuniculus (Rabbit).  
 OC Bkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;  
 OC Mammalia, Butheria, Lagomorpha, Leporidae, Oryctolagus.  
 OC NCBI\_TaxID=9986;  
 RN [1]  
 RP SEQUENCE.  
 RX MEDLINE=95075735; PubMed=7984499;  
 RA Grand D., Schindczek M., Struk K., Shively J., Byssesele V.E.,  
 RA Goebel H., Reeve J.R., Jr.,  
 RT "Characterization of two forms of peptide YY, PYY(1-36) and PYY(3-36),  
 in the rabbit."  
 RL Peptides 15:815-820(1994).

CC -1- SIMILARITY: BELONGS TO THE NPV / PPV / PTV FAMILY.  
 DR HSP, P01303; IRON.  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormone3; 1.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR Annotation.  
 KW SEQUENCE 34 AA; 4024 MW; 02D4B9C38A5FC8D CRC64;  
 SQ  
 Query Match 31.8%; Score 67; DB 6; Length 34;  
 Best Local Similarity 43.3%; Pred. No. 0.045;  
 Matches 13; Conservative 7; Mismatches 10; Indels 0; Gaps 0;  
 QY 4 SQPTPGDPGVEDLIRFYDNLQOMLNCYT 33  
 DB 1 SKRPAGDEASPRELNRYASLRHYINLVIT 30

RESULT 13  
 ID 090WF3 PRELIMINARY; PRT; 99 AA.  
 AC 090WF3;  
 DT 01-DEC-2001 (TREMBLrel. 19, Created)  
 DT 01-DEC-2001 (TREMBLrel. 19, Last sequence update)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last annotation update)  
 DE Peptide YY.  
 GN PYY.  
 OS Paralicthys olivaceus (Flounder).  
 OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 OC Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;  
 OC Acanthomorpha; Acanthopterygii; Perciforma; Pleuronectiformes;  
 OC Pleuronectidae; Paralicthyidae; Paralicthys.  
 OC NCBI\_Taxid=8255;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC TISSUE=Brain;  
 RA Kurikawa T.; Suzuki T.;  
 RT "Development of neuropeptide Y related peptides in the digestive  
 RT organs during the larval stage of Japanese flounder, Paralicthys  
 RT olivaceus";  
 RL Submitted (FEB-2001) to the EMBL/GenBank/DBJ databases.  
 CC -1- SIMILARITY: BELONGS TO THE NPV / PPV / PTV FAMILY.  
 DR EMBL; AB055212; BAB62410.1; -  
 DR InterPro; IPR001955; Pancreatic\_hormn.  
 DR Pfam; PF00159; hormone3; 1.  
 DR PROSITE; PS00276; PANCREATIC\_HORMN.  
 DR ProDom; PD001267; Pancreatic\_hormn; 1.  
 DR SMART; SM00309; PAH; 1.  
 DR PROSITE; PS00265; PANCREATIC\_HORMONE\_1; 1.  
 DR PROSITE; PS50276; PANCREATIC\_HORMONE\_2; 1.  
 DR Annotation.  
 KW SEQUENCE 99 AA; 11179 MW; 32F6C21217CB1984 CRC64;  
 SQ  
 Query Match 30.8%; Score 65; DB 13; Length 99;  
 Best Local Similarity 35.5%; Pred. No. 0.27;  
 Matches 11; Conservative 8; Mismatches 12; Indels 0; Gaps 0;  
 QY 3 PSQPTPGDPGVEDLIRFYDNLQOMLNCYT 33  
 DB 29 PVKPTIPREGATPEADIAKYASLRHYINLVIT 59

RESULT 14  
 ID 08DIF5 PRELIMINARY; PRT; 581 AA.  
 AC 08DIF5;  
 DT 01-MAR-2003 (TREMBLrel. 23, Created)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last sequence update)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last annotation update)  
 DE T11634 protein.  
 DE T11634.

OS Synechococcus elongatus (Thermosynechococcus elongatus).  
 OC Bacteria; Cyanobacteria; Chroococcales; Synechococcus.  
 OC NCBI\_Taxid=32046;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=BP-1;  
 RX MEDLINE=22225144; PubMed=12240834;  
 RA Nakamura Y.; Kaneko T.; Sato S.; Ikeuchi M.; Katoh H.; Sasamoto S.;  
 RA Watanabe A.; Iriyuchi M.; Kawashima K.; Kimura T.; Kishida Y.;  
 RA Kiyokawa C.; Kohara M.; Matsumoto M.; Matsuno A.; Nakazaki N.;  
 RA Shimpō S.; Sugimoto M.; Takeuchi C.; Yamada M.; Tabata S.;  
 RT "Complete genome structure of the thermophilic cyanobacterium  
 RT Thermosynechococcus elongatus BP-1.";  
 RL DNA Res. 9:123-130(2002).  
 DR EMBL; AB005374; BAC09186.1; -  
 KW Complete proteome.  
 SQ SEQUENCE 250 AA; 27453 MW; 429AEDF922699165 CRC64;  
 Query Match 29.1%; Score 63; DB 16; Length 581;  
 Best Local Similarity 46.4%; Pred. No. 3.5;  
 Matches 13; Conservative 0; Mismatches 15; Indels 0; Gaps 0;  
 QY 8 YPGDPGVEDLIRFYDNLQOMLNCYTAA 35  
 DB 245 YPDQDIOGRGLITLYDROQWLEAALAA 272

RESULT 15  
 ID 08DH44 PRELIMINARY; PRT; 250 AA.  
 AC 08DH44;  
 DT 01-MAR-2003 (TREMBLrel. 23, Created)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last sequence update)  
 DT 01-MAR-2003 (TREMBLrel. 23, Last annotation update)  
 DE T112116 protein.  
 GN T112116.  
 OS Synechococcus elongatus (Thermosynechococcus elongatus).  
 OC Bacteria; Cyanobacteria; Chroococcales; Synechococcus.  
 OC NCBI\_Taxid=32046;  
 RN [1]  
 RP SEQUENCE FROM N.A.  
 RC STRAIN=BP-1;  
 RX MEDLINE=22225144; PubMed=12240834;  
 RA Nakamura Y.; Kaneko T.; Sato S.; Ikeuchi M.; Katoh H.; Sasamoto S.;  
 RA Watanabe A.; Iriyuchi M.; Kawashima K.; Kimura T.; Kishida Y.;  
 RA Kiyokawa C.; Kohara M.; Matsumoto M.; Matsuno A.; Nakazaki N.;  
 RA Shimpō S.; Sugimoto M.; Takeuchi C.; Yamada M.; Tabata S.;  
 RT "Complete genome structure of the thermophilic cyanobacterium  
 RT Thermosynechococcus elongatus BP-1.";  
 RL DNA Res. 9:123-130(2002).  
 DR EMBL; AB005376; BAC09668.1; -  
 KW Complete proteome.  
 SQ SEQUENCE 250 AA; 27453 MW; 429AEDF922699165 CRC64;  
 Query Match 29.1%; Score 61.5; DB 16; Length 250;  
 Best Local Similarity 51.7%; Pred. No. 2.2;  
 Matches 15; Conservative 3; Mismatches 10; Indels 1; Gaps 1;  
 QY 2 CPSQPTPGDPGVEDLIRFYDNLQOML 29  
 DB 162 CDSGTFPGDVAVPERLQGFYDHLQVLA 190

Search completed: December 17, 2003, 16:34:34  
 Job time : 37 secs

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OM protein - protein search, using SW model

Run on: December 17, 2003, 16:26:20 / Search time 36 seconds  
(without alignments)  
186.759 Million cell updates/sec

Title: US-10-027-038-11

Perfect score: 211  
Sequence: 1 MCPSPPTYPGDPGVEDLIRFYDNLQOMLNCVTAAAC 36

Scoring table: BLOSUM62  
Gapop 10.0, Gapext 0.5

Searched: 696363 seqs, 186758610 residues

Total number of hits satisfying chosen parameters: 696363

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-Processing: Minimum Match 0%

Maximum Match 100%  
Listing first 45 summaries

Database:

Published Applications AA:  
1: /cgn2\_6/ptodata/1/pubpaa/US07\_PUBCOMB.pep:\*  
2: /cgn2\_6/ptodata/1/pubpaa/PCT\_NEW\_PUB.pep:\*  
3: /cgn2\_6/ptodata/1/pubpaa/US06\_NEW\_PUB.pep:\*  
4: /cgn2\_6/ptodata/1/pubpaa/US06\_PUBCOMB.pep:\*  
5: /cgn2\_6/ptodata/1/pubpaa/US07\_NEW\_PUB.pep:\*  
6: /cgn2\_6/ptodata/1/pubpaa/PCTUS\_PUBCOMB.pep:\*  
7: /cgn2\_6/ptodata/1/pubpaa/US08\_NEW\_PUB.pep:\*  
8: /cgn2\_6/ptodata/1/pubpaa/US08\_PUBCOMB.pep:\*  
9: /cgn2\_6/ptodata/1/pubpaa/US09A\_PUBCOMB.pep:\*  
10: /cgn2\_6/ptodata/1/pubpaa/US09B\_PUBCOMB.pep:\*  
11: /cgn2\_6/ptodata/1/pubpaa/US09C\_PUBCOMB.pep:\*  
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13: /cgn2\_6/ptodata/1/pubpaa/US10A\_PUBCOMB.pep:\*  
14: /cgn2\_6/ptodata/1/pubpaa/US10B\_PUBCOMB.pep:\*  
15: /cgn2\_6/ptodata/1/pubpaa/US10C\_PUBCOMB.pep:\*  
16: /cgn2\_6/ptodata/1/pubpaa/US10\_NEW\_PUB.pep:\*  
17: /cgn2\_6/ptodata/1/pubpaa/US60\_NEW\_PUB.pep:\*  
18: /cgn2\_6/ptodata/1/pubpaa/US60\_PUBCOMB.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	211	100.0	36	US-10-027-038-11	Sequence 11, Appl
2	206	97.6	35	US-10-027-038-14	Sequence 14, Appl
3	193.5	91.7	40	US-10-027-038-22	Sequence 22, Appl
4	193	91.5	41	US-10-027-038-21	Sequence 21, Appl
5	171	81.0	37	US-10-027-038-4	Sequence 4, Appl
6	161	76.3	37	US-10-027-038-3	Sequence 3, Appl
7	156	73.9	36	US-10-027-038-2	Sequence 2, Appl
8	155	73.5	34	US-10-027-038-8	Sequence 8, Appl
9	155	72.5	36	US-10-027-038-5	Sequence 5, Appl
10	153	72.5	36	US-10-027-038-6	Sequence 6, Appl
11	147	69.7	33	US-10-027-038-10	Sequence 10, Appl
12	147	69.7	36	US-10-027-038-1	Sequence 1, Appl
13	147	69.7	37	US-10-027-038-9	Sequence 9, Appl
14	137	64.9	36	US-09-840-085-6	Sequence 6, Appl
15	116	55.0	34	US-09-840-085-54	Sequence 54, Appl

16	111	52.6	34	12	US-09-840-085-56	Sequence 56, Appl
17	110	52.1	31	12	US-09-840-085-59	Sequence 59, Appl
18	110	52.1	34	12	US-09-840-085-55	Sequence 55, Appl
19	107	50.7	34	12	US-09-840-085-53	Sequence 53, Appl
20	106	50.2	41	12	US-09-840-085-49	Sequence 49, Appl
21	101	47.9	31	12	US-09-840-085-38	Sequence 38, Appl
22	98	46.4	39	12	US-09-840-085-8	Sequence 8, Appl
23	96	45.5	31	12	US-09-840-085-32	Sequence 32, Appl
24	96	45.5	41	12	US-09-840-085-47	Sequence 47, Appl
25	95	45.0	39	12	US-09-840-085-9	Sequence 9, Appl
26	93	44.1	31	12	US-09-840-085-60	Sequence 60, Appl
27	91	43.1	50	12	US-09-840-085-64	Sequence 64, Appl
28	90	42.7	39	12	US-09-840-085-10	Sequence 10, Appl
29	90	42.7	39	12	US-09-840-085-11	Sequence 11, Appl
30	90	42.7	39	12	US-09-840-085-12	Sequence 12, Appl
31	90	42.7	39	12	US-09-840-085-14	Sequence 14, Appl
32	90	42.7	43	12	US-09-840-085-62	Sequence 62, Appl
33	90	42.7	50	12	US-09-840-085-63	Sequence 63, Appl
34	89	42.2	41	12	US-09-840-085-52	Sequence 52, Appl
35	88	41.7	43	12	US-09-840-085-61	Sequence 61, Appl
36	86	40.8	36	14	US-10-038-045-5	Sequence 5, Appl
37	84	39.8	36	14	US-10-016-969-4	Sequence 4, Appl
38	84	39.8	97	14	US-10-002-048A-2	Sequence 2, Appl
39	84	39.8	97	14	US-10-036-542-65	Sequence 65, Appl
40	84	39.8	97	15	US-10-236-903-6	Sequence 6, Appl
41	84	39.8	97	15	US-10-205-823-289	Sequence 289, App
42	83	39.3	31	12	US-09-840-085-57	Sequence 57, Appl
43	81	38.4	36	14	US-10-038-045-3	Sequence 3, Appl
44	78	37.0	36	15	US-10-197-954-101	Sequence 101, App
45	77	36.5	34	14	US-10-016-969-5	Sequence 5, Appl

## ALIGNMENTS

RESULT 1  
US-10-027-038-11 ← this case  
Sequence 11, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 11  
LENGTH: 36  
TYPE: PPT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-11

Query Match  
Best Local Similarity 100.0%; Score 211; DB 12;  
Matches 36; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MCPSPPTYPGDPGVEDLIRFYDNLQOMLNCVTAAAC 36  
DB 1 MCPSPPTYPGDPGVEDLIRFYDNLQOMLNCVTAAAC 36

RESULT 2  
US-10-027-038-14  
Sequence 14, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038

CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 14  
LENGTH: 35  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-14

Query Match 97.6%; Score 206; DB 12; Length 35;  
Best Local Similarity 100.0%; Pred. No. 1.8e-19;  
Matches 35; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 CPSQPTPGDPGVEDLIRFYDNLQOMLNCVTAC 36  
DB 1 CPSQPTPGDPGVEDLIRFYDNLQOMLNCVTAC 35

RESULT 3  
US-10-027-038-22

Sequence 22, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 22  
LENGTH: 40  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide-based reagent that combines the SEQ ID  
OTHER INFORMATION: NO:18 interactive domain with the SEQ ID NO:11  
OTHER INFORMATION: peptide backbone.  
US-10-027-038-22

Query Match 91.7%; Score 193.5; DB 12; Length 40;  
Best Local Similarity 87.5%; Pred. No. 8.3e-18;  
Matches 35; Conservative 0; Mismatches 0; Indels 5; Gaps 1;

QY 2 CPSQPTPGDPGVEDLIRFYDNLQOMLNCVTAC 36  
DB 1 CPSQPTPGDPGVEDLIRFYDNLQOMLNCVTAC 40

RESULT 4  
US-10-027-038-21

Sequence 21, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 21  
LENGTH: 41  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide-based reagent that combines the SEQ ID  
OTHER INFORMATION: NO:15 interactive domain with the SEQ ID NO:11  
OTHER INFORMATION: peptide backbone.  
US-10-027-038-21

Query Match 91.5%; Score 193; DB 12; Length 41;  
Best Local Similarity 85.4%; Pred. No. 9.9e-18;  
Matches 35; Conservative 0; Mismatches 0; Indels 6; Gaps 1;

QY 2 CPSQPTPGDPGVEDLIRFYDNLQOMLNCVTAC 36  
DB 1 CPSQPTPGDPGVEDLIRFYDNLQOMLNCVTAC 41

RESULT 5  
US-10-027-038-4

Sequence 4, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 4  
LENGTH: 37  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-4

Query Match 81.0%; Score 171; DB 12; Length 37;  
Best Local Similarity 90.9%; Pred. No. 5.9e-15;  
Matches 30; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 MCSQPTPGDPGVEDLIRFYDNLQOMLNCVT 33  
DB 1 MCSQPTPGDPGVEDLIRFYDNLQOMLNCVT 33

RESULT 6  
US-10-027-038-3

Sequence 3, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 3  
LENGTH: 37  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-3

Query Match 76.3%; Score 161; DB 12; Length 37;  
Best Local Similarity 87.9%; Pred. No. 1.1e-13;  
Matches 29; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 1 MCSQPTPGDPGVEDLIRFYDNLQOMLNCVT 33  
DB 1 MCSQPTPGDPGVEDLIRFYDNLQOMLNCVT 33

RESULT 7  
US-10-027-038-2

Sequence 2, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.

TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 2  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-2

Query Match 73.9%; Score 156; DB 12; Length 36;  
Best Local Similarity 90.3%; Pred. No. 4.9e-13;  
Matches 28; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQQWLNCTV 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQQYLVNVT 32

RESULT 8  
US-10-027-038-8  
Sequence 8, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Oritk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 8  
LENGTH: 34  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-8

Query Match 73.5%; Score 155; DB 12; Length 34;  
Best Local Similarity 87.9%; Pred. No. 6.2e-13;  
Matches 29; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQQWLNCTV 35  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQQYLVNVT 34

RESULT 9  
US-10-027-038-5  
Sequence 5, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Oritk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 5  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-5

Query Match 73.5%; Score 155; DB 12; Length 36;

Best Local Similarity 90.3%; Pred. No. 6.6e-13;  
Matches 28; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQQWLNCTV 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQQYLVNVT 32

RESULT 10  
US-10-027-038-6  
Sequence 6, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Oritk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 6  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-6

Query Match 72.5%; Score 153; DB 12; Length 36;  
Best Local Similarity 90.3%; Pred. No. 1.2e-12;  
Matches 28; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQQWLNCTV 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQQYLVNVT 32

RESULT 11  
US-10-027-038-10  
Sequence 10, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Oritk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027,038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FastSeq for Windows Version 4.0  
SEQ ID NO 10  
LENGTH: 33  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-10

Query Match 69.7%; Score 147; DB 12; Length 33;  
Best Local Similarity 87.1%; Pred. No. 6.4e-12;  
Matches 27; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQQWLNCTV 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQQYLVNVT 32

RESULT 12  
US-10-027-038-1  
Sequence 1, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Oritk, S.  
TITLE OF INVENTION: Modular peptide-based reagent

FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027.038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FASTSEQ for Windows Version 4.0  
SEQ ID NO 1  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Meleagris gallopavo  
US-10-027-038-1

Query Match 69.7%; Score 147; DB 12; Length 36;  
Best Local Similarity 87.1%; Pred. No. 7e-12;  
Matches 27; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQOMLNCVT 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQOYLNVVT 32

RESULT 13  
US-10-027-038-9  
Sequence 9, Application US/10027038  
Publication No. US20030158380A1  
GENERAL INFORMATION:  
APPLICANT: Quirk, S.  
TITLE OF INVENTION: Modular peptide-based reagent  
FILE REFERENCE: 1443.026US1  
CURRENT APPLICATION NUMBER: US/10/027.038  
CURRENT FILING DATE: 2001-12-20  
NUMBER OF SEQ ID NOS: 34  
SOFTWARE: FASTSEQ for Windows Version 4.0  
SEQ ID NO 9  
LENGTH: 37  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: A peptide backbone.  
US-10-027-038-9

Query Match 69.7%; Score 147; DB 12; Length 37;  
Best Local Similarity 87.1%; Pred. No. 7.2e-12;  
Matches 27; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQOMLNCVT 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQOYLNVVT 32

RESULT 14  
US-09-840-085-6  
Sequence 6, Application US/09840085  
Publication No. US20030166240A1  
GENERAL INFORMATION:  
APPLICANT: Schepartz Shrader, Alanna  
APPLICANT: Chin, Jason W. K.  
APPLICANT: Zutschi, Reena  
APPLICANT: Rutledge, Stacey E.  
APPLICANT: Kehlbeck Martin, Joanne D.  
TITLE OF INVENTION: DNA and Protein Binding Miniature Proteins  
FILE REFERENCE: 44574-5099-US  
CURRENT APPLICATION NUMBER: US/09/840.085  
CURRENT FILING DATE: 2001-04-24  
PRIOR APPLICATION NUMBER: US 60/199,408  
PRIOR FILING DATE: 2000-04-24  
PRIOR APPLICATION NUMBER: US 60/240,566  
PRIOR FILING DATE: 2000-10-13  
PRIOR APPLICATION NUMBER: US PROVISIONAL  
PRIOR FILING DATE: 2001-01-13  
PRIOR APPLICATION NUMBER: US PROVISIONAL  
PRIOR FILING DATE: 2001-02-23  
NUMBER OF SEQ ID NOS: 73

SOFTWARE: PatentIn Ver. 2.1  
SEQ ID NO 6  
LENGTH: 36  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: Description of Artificial Sequence: Segment of  
US-09-840-085-6

Query Match 64.9%; Score 137; DB 12; Length 36;  
Best Local Similarity 80.6%; Pred. No. 1.3e-10;  
Matches 25; Conservative 3; Mismatches 3; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQOMLNCVT 33  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQOYLNVVT 32

RESULT 15  
US-09-840-085-54  
Sequence 54, Application US/09840085  
Publication No. US20030166240A1  
GENERAL INFORMATION:  
APPLICANT: Schepartz Shrader, Alanna  
APPLICANT: Chin, Jason W. K.  
APPLICANT: Zutschi, Reena  
APPLICANT: Rutledge, Stacey E.  
APPLICANT: Kehlbeck Martin, Joanne D.  
TITLE OF INVENTION: DNA and Protein Binding Miniature Proteins  
FILE REFERENCE: 44574-5099-US  
CURRENT APPLICATION NUMBER: US/09/840.085  
CURRENT FILING DATE: 2001-04-24  
PRIOR APPLICATION NUMBER: US 60/199,408  
PRIOR FILING DATE: 2000-04-24  
PRIOR APPLICATION NUMBER: US 60/240,566  
PRIOR FILING DATE: 2000-10-13  
PRIOR APPLICATION NUMBER: US PROVISIONAL  
PRIOR FILING DATE: 2001-01-13  
PRIOR APPLICATION NUMBER: US PROVISIONAL  
PRIOR FILING DATE: 2001-02-23  
NUMBER OF SEQ ID NOS: 73  
SOFTWARE: PatentIn Ver. 2.1  
SEQ ID NO 54  
LENGTH: 34  
TYPE: PRT  
ORGANISM: Artificial Sequence  
FEATURE:  
OTHER INFORMATION: Description of Artificial Sequence: PMW02, MyoD  
US-09-840-085-54

Query Match 55.0%; Score 116; DB 12; Length 34;  
Best Local Similarity 70.0%; Pred. No. 6.3e-08;  
Matches 21; Conservative 3; Mismatches 6; Indels 0; Gaps 0;

QY 3 PSQPTPGDGPVEDLIRFYDNLQOMLNCVT 32  
DB 2 PSQPTPGDDAPVEDLIRFYDNLQOYLNVVT 31

Search completed: December 17, 2003, 16:32:26  
Job time : 37 secs